ARRICK, JAMES W. TRADE

## SEQUENCE LISTING

WYCOFF, KEITH L.

<120> NOVEL IMMUNOADHESINS FOR TREATING AND PREVENTING VIRAL AND BACTERIAL DISEASES

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<151> 2001-04-28

<150> 60/200,298

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<170> PatentIn Ver. 2.1

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Leu Arg Val Ala Ala Glu Asp Trp Lys Lys Gly Glu Thr Phe Ser Cys 740 745 750

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Lys Ser Val Thr Cys His Val Lys His Tyr Thr Asn Pro Ser Gln Asp

Val Thr Val Pro Cys Pro Val Pro Ser Thr Pro Pro Thr Pro Ser Pro 105 110 100

Ser Thr Pro Pro Thr Pro Ser Pro Ser Cys Cys His Pro Arg Leu Ser 115 120 Leu His Arg Pro Ala Leu Glu Asp Leu Leu Gly Ser Glu Ala Asn Leu Thr Cys Thr Leu Thr Gly Leu Arg Asp Ala Ser Gly Val Thr Phe 150 Thr Trp Thr Pro Ser Ser Gly Lys Ser Ala Val Gln Gly Pro Pro Glu 170 Arg Asp Leu Cys Gly Cys Tyr Ser Val Ser Ser Val Leu Pro Gly Cys 185 Ala Glu Pro Trp Asn His Gly Lys Thr Phe Thr Cys Thr Ala Ala Tyr Pro Glu Ser Lys Thr Pro Leu Thr Ala Thr Leu Ser Lys Ser Gly Asn Thr Phe Arg Pro Glu Val His Leu Leu Pro Pro Pro Ser Glu Glu Leu 225 230 Ala Leu Asn Glu Leu Val Thr Leu Thr Cys Leu Ala Arg Gly Phe, Ser 245 Pro Lys Asp Val Leu Val Arg Trp Leu Gln Gly Ser Gln Glu Leu Pro 265 Arg Glu Lys Tyr Leu Thr Trp Ala Ser Arg Gln Glu Pro Ser Gln Gly Thr Thr Thr Phe Ala Val Thr Ser Ile Leu Arg Val Ala Ala Glu Asp Trp Lys Lys Gly Asp Thr Phe Ser Cys Met Val Gly His Glu Ala Leu Pro Leu Ala Phe Thr Gln Lys Thr Ile Asp Arg Leu Ala Gly Lys Pro Thr His Val Asn Val Ser Val Val Met Ala Glu Val Asp Gly Thr Cys

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Thr Ala Arg Asn Phe Pro Pro Ser Gln Asp Ala Ser Gly Asp Leu Tyr 50 55 60

Thr Thr Ser Ser Gln Leu Thr Leu Pro Ala Thr Gln Cys Pro Asp Gly 65 70 75 80

Lys Ser Val Thr Cys His Val Lys His Tyr Thr Asn Pro Ser Gln Asp 85 90 95

Val Thr Val Pro Cys Pro Val Pro Pro Pro Pro Cys Cys His Pro
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Arg Leu Ser Leu His Arg Pro Ala Leu Glu Asp Leu Leu Gly Ser 115 120 125

Glu Ala Asn Leu Thr Cys Thr Leu Thr Gly Leu Arg Asp Ala Ser Gly 130 135 140

Ala Thr Phe Thr Trp Thr Pro Ser Ser Gly Lys Ser Ala Val Gln Gly 145 150 155 160

Pro Pro Glu Arg Asp Leu Cys Gly Cys Tyr Ser Val Ser Ser Val Leu 165 170 175

Pro Gly Cys Ala Gln Pro Trp Asn His Gly Glu Thr Phe Thr Cys Thr 180 185 190 Ala Ala His Pro Glu Leu Lys Thr Pro Leu Thr Ala Asn Ile Thr Lys 200 Ser Gly Asn Thr Phe Arg Pro Glu Val His Leu Leu Pro Pro Pro Ser Glu Glu Leu Ala Leu Asn Glu Leu Val Thr Leu Thr Cys Leu Ala Arg 235 230 Gly Phe Ser Pro Lys Asp Val Leu Val Arg Trp Leu Gln Gly Ser Gln 250 245 Glu Leu Pro Arg Glu Lys Tyr Leu Thr Trp Ala Ser Arg Gln Glu Pro 265 Ser Gln Gly Thr Thr Thr Phe Ala Val Thr Ser Ile Leu Arg Val Ala 275 Ala Glu Asp Trp Lys Lys Gly Asp Thr Phe Ser Cys Met Val Gly His Glu Ala Leu Pro Leu Ala Phe Thr Gln Lys Thr Ile Asp Arg Leu Ala 315 305 310 Gly Lys Pro Thr His Val Asn Val Ser Val Val Met Ala Glu Val Asp 330 325 Gly Thr Cys Tyr 340

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Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser 50 55 60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr 65 70 75 80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys 85 90 95

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Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro 115 120 125

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Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp 145 150 155 160

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His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn 195 200 205

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Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu 225 230 235 240

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Thr Val Glu Arg Lys Cys Cys Val Glu Cys Pro Pro Cys Pro Ala Pro 115

Pro Val Ala Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp 125

Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp 130 135 140

Val Ser His Glu Asp Pro Glu Val Gln Phe Asn Trp Tyr Val Asp Gly
145 150 155 160

Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Phe Asn 165 170 175

Ser Thr Phe Arg Val Val Ser Val Leu Thr Val Val His Gln Asp Trp 180 185 190

Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Gly Leu Pro 195 200 205

Ala Pro Ile Glu Lys Thr Ile Ser Lys Thr Lys Gly Gln Pro Arg Glu 210 215 220

Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn 225 230 235 240

Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile 245 250 255

Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr 260 265 270

Thr Pro Pro Met Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys 275 280 285

Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys 290 295 300

Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu 305 310 315 320

Ser Leu Ser Pro Gly Lys 325

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<211> 1134

<212> DNA

<213> Homo sapiens

<400> 23

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					caaatcttgt	
					acctccccca	
					aaggtgccca	
					caaggatacc	
					ccacgaagac	
					caagacaaag	
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	accgcttcac					1134

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<211> 377

<212> PRT

<213> Homo sapiens

<400> 24

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser Arg
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Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr 20 25 30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser 35 40 45

Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser 50 55 60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr 65 70 75 80

Tyr Thr Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
85 90 95

Arg Val Glu Leu Lys Thr Pro Leu Gly Asp Thr Thr His Thr Cys Pro 100 105 110

Arg Cys Pro Glu Pro Lys Ser Cys Asp Thr Pro Pro Pro Cys Pro Arg 115 120 125

Cys Pro Glu Pro Lys Ser Cys Asp Thr Pro Pro Pro Cys Pro Arg Cys 130 135 140

Pro Glu Pro Lys Ser Cys Asp Thr Pro Pro Pro Cys Pro Arg Cys Pro 145 150 155 160

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys 165 170 175 Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val 185 Val Val Asp Val Ser His Glu Asp Pro Glu Val Gln Phe Lys Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu 215 Gln Tyr Asn Ser Thr Phe Arg Val Val Ser Val Leu Thr Val Leu His 230 235 Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Thr Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met 280 275 Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro 295 Ser Asp Ile Ala Val Glu Trp Glu Ser Ser Gly Gln Pro Glu Asn Asn 310 Tyr Asn Thr Thr Pro Pro Met Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Ile 345 Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn Arg Phe Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys 375

<210> 25 <211> 984 <212> DNA <213> Homo sapiens

<400> 25

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aaccaggtca gcctgacctg cctggtcaaa ggcttctacc ccagcgacat cgccgtggag 780 tgggagagca atgggcagcc ggagaacaac tacaagacca cgcctcccgt gctggactcc 840 gacggctcct tcttcctcta cagcaggcta accgtggaca agagcaggtg gcaggagggg 900 aatgtcttct catgctccgt gatgcatgag gctctgcaca accactacac acagaagagc 960 ctctccctgt ctctgggtaa atga

<210> 26

<211> 327

<212> PRT

<213> Homo sapiens

<400> 26

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser Arg

Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
20 25 30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser 35 40 45

Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser 50 55 60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Lys Thr 65 70 75 80

Tyr Thr Cys Asn Val Asp His Lys Pro Ser Asn Thr Lys Val Asp Lys
85 90 95

Arg Val Glu Ser Lys Tyr Gly Pro Pro Cys Pro Ser Cys Pro Ala Pro 100 105 110

Glu Phe Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys 115 120 125

Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val 130 135 140

Asp Val Ser Gln Glu Asp Pro Glu Val Gln Phe Asn Trp Tyr Val Asp 145 150 155 160

Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Phe 165 170 175

Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp 180 185 190

Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Gly Leu 195 200 205

Pro Ser Ser Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg 210 215 220

Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Gln Glu Glu Met Thr Lys 225 230 235 240 Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp 245 250 255

Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys 260 265 270

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser 275 280 285

Arg Leu Thr Val Asp Lys Ser Arg Trp Gln Glu Gly Asn Val Phe Ser 290 295 300

Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser 305 310 315 320

Leu Ser Leu Ser Leu Gly Lys 325

<210> 27

<211> 300

<212> DNA

<213> Homo sapiens

<400> 27

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<210> 28

<211> 383

<212> PRT

<213> Homo sapiens

<400> 28

Ala Pro Thr Lys Ala Pro Asp Val Phe Pro Ile Ile Ser Gly Cys Arg 1 5 10 15

His Pro Lys Asp Asn Ser Pro Val Val Leu Ala Cys Leu Ile Thr Gly 20 25 30

Tyr His Pro Thr Ser Val Thr Val Thr Trp Tyr Met Gly Thr Gln Ser 35 40 45

Gln Pro Gln Arg Thr Phe Pro Glu Ile Gln Arg Arg Asp Ser Tyr Tyr 50 55 60

Met Thr Ser Ser Gln Leu Ser Thr Pro Leu Gln Gln Trp Arg Gln Gly 65 70 75 80

Glu Tyr Lys Cys Val Val Gln His Thr Ala Ser Lys Ser Lys Glu 85 90 95

Ile Phe Arg Trp Pro Glu Ser Pro Lys Ala Gln Ala Ser Ser Val Pro 100 105 110 Thr Ala Gln Pro Gln Ala Glu Gly Ser Leu Ala Lys Ala Thr Thr Ala 115 120 125

Pro Ala Thr Thr Arg Asn Thr Gly Arg Gly Glu Glu Lys Lys Lys 130 135 140

Glu Lys Glu Lys Glu Glu Glu Glu Glu Arg Glu Thr Lys Thr Pro Glu 145 150 155 160

Cys Pro Ser His Thr Gln Pro Leu Gly Val Tyr Leu Leu Thr Pro Ala 165 170 175

Val Gln Asp Leu Trp Leu Arg Asp Lys Ala Thr Phe Thr Cys Phe Val 180 185 190

Val Gly Ser Asp Leu Lys Asp Ala His Leu Thr Trp Glu Val Ala Gly
195 200 205

Lys Val Pro Thr Gly Gly Val Glu Glu Gly Leu Leu Glu Arg His Ser 210 215 220

Asn Gly Ser Gln Ser Gln His Ser Arg Leu Thr Leu Pro Arg Ser Leu 225 230 235 240

Trp Asn Ala Gly Thr Ser Val Thr Cys Thr Leu Asn His Pro Ser Leu 245 250 255

Pro Pro Gln Arg Leu Met Ala Leu Arg Glu Pro Ala Ala Gln Ala Pro 260 265 270

Val Lys Leu Ser Leu Asn Leu Leu Ala Ser Ser Asp Pro Pro Glu Ala 275 280 285

Ala Ser Trp Leu Leu Cys Glu Val Ser Gly Phe Ser Pro Pro Asn Ile 290 295 300

Leu Leu Met Trp Leu Glu Asp Gln Arg Glu Val Asn Thr Ser Gly Phe 305 310 315 320

Ala Pro Ala Arg Pro Pro Pro Gln Pro Gly Ser Thr Thr Phe Trp Ala
325 330 335

Trp Ser Val Leu Arg Val Pro Ala Pro Pro Ser Pro Gln Pro Ala Thr 340 345 350

Tyr Thr Cys Val Val Ser His Glu Asp Ser Arg Thr Leu Leu Asn Ala 355 360 365

Ser Arg Ser Leu Glu Val Ser Tyr Val Thr Asp His Gly Pro Met 370 375 380

<210> 29

<211> 300

<212> DNA

<213> Homo sapiens

<400> 29

<210> 30

<211> 429

<212> PRT

<213> Homo sapiens

<400> 30

Pro Thr Lys Ala Pro Asp Val Phe Pro Ile Ile Ser Gly Cys Arg His 1 5 10 15

Pro Lys Asp Asn Ser Pro Val Val Leu Ala Cys Leu Ile Thr Gly Tyr
20 25 30

His Pro Thr Ser Val Thr Val Thr Trp Tyr Met Gly Thr Gln Ser Gln 35 40 45

Pro Gln Arg Thr Phe Pro Glu Ile Gln Arg Arg Asp Ser Tyr Tyr Met 50 55 60

Thr Ser Ser Gln Leu Ser Thr Pro Leu Gln Gln Trp Arg Gln Gly Glu 65 70 75 80

Tyr Lys Cys Val Val Gln His Thr Ala Ser Lys Ser Lys Lys Glu Ile 85 90 95

Phe Arg Trp Pro Glu Ser Pro Lys Ala Gln Ala Ser Ser Val Pro Thr 100 105 110

Ala Gln Pro Gln Ala Glu Gly Ser Leu Ala Lys Ala Thr Thr Ala Pro 115 120 125

Ala Thr Thr Arg Asn Thr Gly Arg Gly Glu Glu Lys Lys Lys Glu
130 135 140

Lys Glu Lys Glu Glu Glu Glu Glu Glu Arg Glu Thr Lys Thr Pro Glu Cys 145 150 155 160

Pro Ser His Thr Gln Pro Leu Gly Val Tyr Leu Leu Thr Pro Ala Val

Gln Asp Leu Trp Leu Arg Asp Lys Ala Thr Phe Thr Cys Phe Val Val 180 185 190

Gly Ser Asp Leu Lys Asp Ala His Leu Thr Trp Glu Val Ala Gly Lys 195 200 205

Val Pro Thr Gly Gly Val Glu Glu Gly Leu Leu Glu Arg His Ser Asn 210 215 220

Gly Ser Gln Ser Gln His Ser Arg Leu Thr Leu Pro Arg Ser Leu Trp 225 230 235 240 Asn Ala Gly Thr Ser Val Thr Cys Thr Leu Asn His Pro Ser Leu Pro Pro Gln Arg Leu Met Ala Leu Arg Glu Pro Ala Ala Gln Ala Pro Val 260 Lys Leu Ser Leu Asn Leu Leu Ala Ser Ser Asp Pro Pro Glu Ala Ala 280 Ser Trp Leu Leu Cys Glu Val Ser Gly Phe Ser Pro Pro Asn Ile Leu Leu Met Trp Leu Glu Asp Gln Arg Glu Val Asn Thr Ser Gly Phe Ala Pro Ala Arg Pro Pro Pro Gln Pro Arg Ser Thr Thr Phe Trp Ala Trp 325 330 Ser Val Leu Arg Val Pro Ala Pro Pro Ser Pro Gln Pro Ala Thr Tyr 345 Thr Cys Val Val Ser His Glu Asp Ser Arg Thr Leu Leu Asn Ala Ser 360 Arg Ser Leu Glu Val Ser Tyr Leu Ala Met Thr Pro Leu Ile Pro Gln 375 Ser Lys Asp Glu Asn Ser Asp Asp Tyr Thr Thr Phe Asp Asp Val Gly 400 Ser Leu Trp Thr Thr Leu Ser Thr Phe Val Ala Leu Phe Ile Leu Thr 410 405 Leu Leu Tyr Ser Gly Ile Val Thr Phe Ile Lys Val Lys 425 <210> 31 <211> 500 <212> DNA <213> Homo sapiens <400> 31 gaagctgggg agaggagagc acagtggtta agtcagtccc tgcagcccaa ctgctcccga 60 aggtccggcc acagctgctc tcgtttgctc tcccctgcag agtgtccgag ccacacccag 120 cctcttqqcq tctacctgct aacccctgca gtgcaggacc tgtggctccg ggacaaagcc 180 accttcacct gcttcgtggt gggcagtgac ctgaaggatg ctcacctgac ctgggaggtg 240 gctgggaagg tccccacagg gggcgtggag gaagggctgc tggagcggca cagcaacggc 300 teccagagee ageacageeg tetgaceetg eccaggteet tgtggaaege ggggaeetee 360 gtcacctgca cactgaacca tcccagcctc ccaccccaga ggttgatggc gctgagagaa 420

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<212> PRT

<213> Homo sapiens

<400> 32

Pro Thr Lys Ala Pro Asp Val Phe Pro Ile Ile Ser Gly Cys Arg His 1 5 10 15

Pro Lys Asp Asn Ser Pro Val Val Leu Ala Cys Leu Ile Thr Gly Tyr
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His Pro Thr Ser Val Thr Val Thr Trp Tyr Met Gly Thr Gln Ser Gln 35 40 45

Pro Gln Arg Thr Phe Pro Glu Ile Gln Arg Arg Asp Ser Tyr Tyr Met 50 55 60

Thr Ser Ser Gln Leu Ser Thr Pro Leu Gln Gln Trp Arg Gln Gly Glu 65 70 75 80

Tyr Lys Cys Val Val Gln His Thr Ala Ser Lys Ser Lys Glu Ile 85 90 95

Phe Arg Trp Pro Glu Ser Pro Lys Ala Gln Ala Ser Ser Val Pro Thr
100 105 110

Ala Gln Pro Gln Ala Glu Gly Ser Leu Ala Lys Ala Thr Thr Ala Pro 115 120 125

Ala Thr Thr Arg Asn Thr Gly Arg Gly Glu Glu Lys Lys Lys Glu 130 135 140

Lys Glu Lys Glu Glu Glu Glu Glu Glu Arg Glu Thr Lys Thr Pro Glu Cys 145 150 155 160

Pro Ser His Thr Gln Pro Leu Gly Val Tyr Leu Leu Thr Pro Ala Val 165 170 175

Gln Asp Leu Trp Leu Arg Asp Lys Ala Thr Phe Thr Cys Phe Val Val 180 185 190

Gly Ser Asp Leu Lys Asp Ala His Leu Thr Trp Glu Val Ala Gly Lys 195 200 205

Val Pro Thr Gly Gly Val Glu Glu Gly Leu Leu Glu Arg His Ser Asn 210 215 220

Gly Ser Gln Ser Gln His Ser Arg Leu Thr Leu Pro Arg Ser Leu Trp 225 230 235 240

Asn Ala Gly Thr Ser Val Thr Cys Thr Leu Asn His Pro Ser Leu Pro 245 250 255

Pro Gln Arg Leu Met Ala Leu Arg Glu Pro Ala Ala Gln Ala Pro Val 260 265 270

Lys Leu Ser Leu Asn Leu Leu Ala Ser Ser Asp Pro Pro Glu Ala Ala 275 280 285

31 Ser Trp Leu Cys Glu Val Ser Gly Phe Ser Pro Pro Asn Ile Leu 295 Leu Met Trp Leu Glu Asp Gln Arg Glu Val Asn Thr Ser Gly Phe Ala 315 310 Pro Ala Arg Pro Pro Pro Gln Pro Arg Ser Thr Thr Phe Trp Ala Trp 330 325 Ser Val Leu Arg Val Pro Ala Pro Pro Ser Pro Gln Pro Ala Thr Tyr 345 Thr Cys Val Val Ser His Glu Asp Ser Arg Thr Leu Leu Asn Ala Ser 355 Arg Ser Leu Glu Val Ser Tyr Val Thr Asp His Gly Pro Met Lys 375 <210> 33 <211> 500 <212> DNA <213> Homo sapiens <400> 33 ccacaggaaa ggagaaggga ggcaccacac cctggccggc cccacttctc tcccagtgcc 60 cccgtggcca gagcctgaca gccccccac ctccccgcag ctgcgcaggc acccgtcaag 120 ctttctctga acctgctggc ctcgtctgac cctcccgagg cggcctcgtg gctcctgtgt 180 gaggtgtctg gcttctcgcc ccccaacatc ctcctgatgt ggctggagga ccagcgtgag 240 gtgaacactt ctgggtttgc ccccgcacgc cccctccac agcccaggag caccacgttc 300 tgggcctgga gtgtgctgcg tgtcccagcc ccgcccagcc ctcagccagc cacctacacg 360 tgtgtggtca gccacgagga ctcccggact ctgctcaacg ccagccggag cctagaagtc 420 agctgtgagt cacccccagg ccagggttgg gacggggact ctgagggggg ccataaggag 480 500 ctggaatcca tactaggcag <210> 34 <400> 34 000 <210> 35 <211> 26 <212> PRT <213> Homo sapiens Pro Thr Lys Ala Pro Asp Val Phe Pro Ile Ile Ser Gly Cys Arg His Pro Lys Asp Asn Ser Pro Val Val Leu Ala

<210> 36 <211> 100 <212> DNA

<213> Homo sapiens

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<212> DNA
<213> Homo sapiens
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ccacgtttgt ggccctcttc atcctcaccc tcctctacag cggcattgtc actttcatca 180
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<210> 45
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<213> Homo sapiens
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<400> 45

Met Asp Trp Thr Trp Ile Leu Phe Leu Val Ala Ala Ala Thr Arg Val
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His Ser Gln Thr Gln Leu Val Gln Ser Gly Ala Glu Val Arg Lys Pro 20 25 30

Gly Ala Ser Val Arg Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Ile 35 40 45

Asp Ser Tyr Ile His Trp Ile Arg Gln Ala Pro Gly His Gly Leu Glu
50 60

Trp Val Gly Trp Ile Asn Pro Asn Ser Gly Gly Thr Asn Tyr Ala Pro 65 70 75 80

Arg Phe Gln Gly Arg Val Thr Met Thr Arg Asp Ala Ser Phe Ser Thr 85 90 95

Ala Tyr Met Asp Leu Arg Ser Leu Arg Ser Asp Asp Ser Ala Val Phe 100 105 110

Tyr Cys Ala Lys Ser Asp Pro Phe Trp Ser Asp Tyr Tyr Asn Phe Asp 115 120 125

Tyr Ser Tyr Thr Leu Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val 130 135 140

Ser Ser Ala Ser Thr Gln Ser Pro Ser Val Phe Pro Leu Thr Arg Cys 145 150 155 160

Cys Lys Asn Ile Pro Ser Asn Ala Thr Ser Val Thr Leu Gly Cys Leu 165 170 175

Ala Thr Gly Tyr Phe Pro Glu Pro Val Met Val Thr Trp Asp Thr Gly 180 185 190

Ser Leu Asn Gly Thr Thr Met Thr Leu Pro Ala Thr Thr Leu Thr Leu 195 200 205

Ser Gly His Tyr Ala Thr Ile Ser Leu Leu Thr Val Ser Gly Ala Trp 210 215 220

Ala Lys Gln Met Phe Thr Cys Arg Val Ala His Thr Pro Ser Ser Thr 225 230 235 240

Asp Trp Val Asp Asn Lys Thr Phe Ser Val Cys Ser Arg Asp Phe Thr 245 250 255

Pro Pro Thr Val Lys Ile Leu Gln Ser Ser Cys Asp Gly Gly His

Phe Pro Pro Thr Ile Gln Leu Cys Leu Val Ser Gly Tyr Thr Pro 275 280 285

Gly Thr Ile Asn Ile Thr Trp Leu Glu Asp Gly Gln Val Met Asp Val 290 295 300 Asp Leu Ser Thr Ala Ser Thr Thr Gln Glu Gly Glu Leu Ala Ser Thr Gln Ser Glu Leu Thr Leu Ser Gln Lys His Trp Leu Ser Asp Arg Thr 325 330 Tyr Thr Cys Gln Val Thr Tyr Gln Gly His Thr Phe Glu Asp Ser Thr Lys Lys Cys Ala Asp Ser Asn Pro Arg Gly Val Ser Ala Tyr Leu Ser 360 Arg Pro Ser Pro Phe Asp Leu Phe Ile Arg Lys Ser Pro Thr Ile Thr Cys Leu Val Val Asp Leu Ala Pro Ser Lys Gly Thr Val Asn Leu Thr 390 Trp Ser Arg Ala Ser Gly Lys Pro Val Asn His Ser Thr Arg Lys Glu 405 410 Glu Lys Gln Arg Asn Gly Thr Leu Thr Val Thr Ser Thr Leu Pro Val 425 Gly Thr Arq Asp Trp Ile Glu Gly Glu Thr Tyr Gln Cys Arg Val Thr His Pro His Leu Pro Arg Ala Leu Met Arg Ser Thr Thr Lys Thr Ser Gly Pro Arg Ala Ala Pro Glu Val Tyr Ala Phe Ala Thr Pro Glu Trp 475 Pro Gly Ser Arg Asp Lys Arg Thr Leu Ala Cys Leu Ile Gln Asn Phe Met Pro Glu Asp Ile Ser Val Gln Trp Leu His Asn Glu Val Gln Leu 505 Pro Asp Ala Arg His Ser Thr Thr Gln Pro Arg Lys Thr Lys Gly Ser Gly Phe Phe Val Phe Ser Arg Leu Glu Val Thr Arg Ala Glu Trp Glu 535 Gln Lys Asp Glu Phe Ile Cys Arg Ala Val His Glu Ala Ala Ser Pro 545

Ser Gln Thr Val Gln Arg Ala Val Ser Val Asn Pro Gly Lys

565

570

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gcaaactacg cacagaagtt ccagggcaga gtcacgatta ccgcggacga atccacgagc 360
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aaaaccggga tcctggggcc gtatagcagt ggctggtacc cgaactcgga ctactactac 480
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                             40
 Ser Ser Tyr Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
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55

Glu Trp Met Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala 75 Gln Lys Phe Gln Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala Lys Thr Gly Ile Leu Gly Pro Tyr Ser Ser Gly Trp 120 Tyr Pro Asn Ser Asp Tyr Tyr Tyr Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Ser Ala Ser Ala Pro Thr Leu Phe Pro Leu Val Ser Cys Glu Asn Ser Pro Ser Asp Thr Ser Ser Val 165 Ala Val Gly Cys Leu Ala Gln Asp Phe Leu Pro Asp Ser Ile Thr Phe 185 Ser Trp Lys Tyr Lys Asn Asn Ser Asp Ile Ser Ser Thr Arg Gly Phe 195 200 Pro Ser Val Leu Arg Gly Gly Lys Tyr Ala Ala Thr Ser Gln Val Leu Leu Pro Ser Lys Asp Val Met Gln Gly Thr Asp Glu His Val Val Cys Lys Val Gln His Pro Asn Gly Asn Lys Glu Lys Asn Val Pro Leu Pro Val Ile Ala Glu Leu Pro Pro Lys Val Ser Val Phe Val Pro Pro Arg Asp Gly Phe Phe Gly Asn Pro Arg Ser Lys Ser Lys Leu Ile Cys Gln Ala Thr Gly Phe Ser Pro Arg Gln Ile Gln Val Ser Trp Leu Arg Glu Gly Lys Gln Val Gly Ser Gly Val Thr Thr Asp Gln Val Gln Ala Glu Ala Lys Glu Ser Gly Pro Thr Thr Tyr Lys Val Thr Ser Thr Leu Thr 330 325 Ile Lys Glu Ser Asp Trp Leu Ser Gln Ser Met Phe Thr Cys Arg Val 340 Asp His Arg Gly Leu Thr Phe Gln Gln Asn Ala Ser Ser Met Cys Val

Pro Asp Gln Asp Thr Ala Ile Arg Val Phe Ala Ile Pro Pro Ser Phe 370 375 380

Ala Ser Ile Phe Leu Thr Lys Ser Thr Lys Leu Thr Cys Leu Val Thr 385 390 395 400

Asp Leu Thr Thr Tyr Asp Ser Val Thr Ile Ser Trp Thr Arg Gln Asn 405 410 415

Gly Glu Ala Val Lys Thr His Thr Asn Ile Ser Glu Ser His Pro Asn 420 425 430

Ala Thr Phe Ser Ala Val Gly Glu Ala Ser Ile Cys Glu Asp Asp Trp
435 440 445

Asn Ser Gly Glu Arg Phe Thr Cys Thr Val Thr His Thr Asp Leu Pro 450 455 460

Ser Pro Leu Lys Gln Thr Ile Ser Arg Pro Lys Gly Val Ala Leu His 465 470 475 480

Arg Pro Asp Val Tyr Leu Leu Pro Pro Ala Arg Glu Gln Leu Asn Leu 485 490 495

Arg Glu Ser Ala Thr Ile Thr Cys Leu Val Thr Gly Phe Ser Pro Ala 500 505 510

Asp Val Phe Val Gln Trp Met Gln Arg Gly Gln Pro Leu Ser Pro Glu 515 520 525

Lys Tyr Val Thr Ser Ala Pro Met Pro Glu Pro Gln Ala Pro Gly Arg 530 535 540

Tyr Phe Ala His Ser Ile Leu Thr Val Ser Glu Glu Glu Trp Asn Thr 545 550 555 560

Gly Glu Thr Tyr Thr Cys Val Val Ala His Glu Ala Leu Pro Asn Arg 565 570 575

Val Thr Glu Arg Thr Val Asp Lys Ser Thr Glu Gly Glu Val Ser Ala 580 585 590

Asp Glu Glu Gly Phe Glu Asn Leu Trp Ala Thr Ala Ser Thr Phe Ile 595 600 605

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Lys Val Lys 625

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<211> 822

<212> PRT

<213> Artificial Sequence

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 encoded by plasmid pSSPICAMHuA2

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Ile Leu Pro Arg Gly Gly Ser Val Leu Val Thr Cys Ser Thr Ser Cys 35 40 45

Asp Gln Pro Lys Leu Leu Gly Ile Glu Thr Pro Leu Pro Lys Lys Glu 50 55 60

Leu Leu Pro Gly Asn Asn Arg Lys Val Tyr Glu Leu Ser Asn Val 65 70 75 80

Gln Glu Asp Ser Gln Pro Met Cys Tyr Ser Asn Cys Pro Asp Gly Gln 85 90 95

Ser Thr Ala Lys Thr Phe Leu Thr Val Tyr Trp Thr Pro Glu Arg Val

Glu Leu Ala Pro Leu Pro Ser Trp Gln Pro Val Gly Lys Asn Leu Thr 115 120 125

Leu Arg Cys Gln Val Glu Gly Gly Ala Pro Arg Ala Asn Leu Thr Val 130 135 140

Val Leu Leu Arg Gly Glu Lys Glu Leu Lys Arg Glu Pro Ala Val Gly
145 150 155 160

Glu Pro Ala Glu Val Thr Thr Thr Val Leu Val Arg Arg Asp His His
165 170 175

Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu Asp Leu Arg Pro Gln Gly
180 185 190

Leu Glu Leu Phe Glu Asn Thr Ser Ala Pro Tyr Gln Leu Gln Thr Phe
195 200 205

Val Leu Pro Ala Thr Pro Pro Gln Leu Val Ser Pro Arg Val Leu Glu 210 215 220

Val Asp Thr Gln Gly Thr Val Val Cys Ser Leu Asp Gly Leu Phe Pro 225 230 235 240

Val Ser Glu Ala Gln Val His Leu Ala Leu Gly Asp Gln Arg Leu Asn
245
250
255

Pro Thr Val Thr Tyr Gly Asn Asp Ser Phe Ser Ala Lys Ala Ser Val 260 265 270

Ser Val Thr Ala Glu Asp Glu Gly Thr Gln Arg Leu Thr Cys Ala Val 275 280 285 Ile Leu Gly Asn Gln Ser Gln Glu Thr Leu Gln Thr Val Thr Ile Tyr Ser Phe Pro Ala Pro Asn Val Ile Leu Thr Lys Pro Glu Val Ser Glu Gly Thr Glu Val Thr Val Lys Cys Glu Ala His Pro Arg Ala Lys Val 330 325 Thr Leu Asn Gly Val Pro Ala Gln Pro Leu Gly Pro Arg Ala Gln Leu Leu Leu Lys Ala Thr Pro Glu Asp Asn Gly Arg Ser Phe Ser Cys Ser 360 Ala Thr Leu Glu Val Ala Gly Gln Leu Ile His Lys Asn Gln Thr Arg Glu Leu Arg Val Leu Tyr Gly Pro Arg Leu Asp Glu Arg Asp Cys Pro Gly Asn Trp Thr Trp Pro Glu Asn Ser Gln Gln Thr Pro Met Cys Gln 405 Ala Trp Gly Asn Pro Leu Pro Glu Leu Lys Cys Leu Lys Asp Gly Thr Phe Pro Leu Pro Ile Gly Glu Ser Val Thr Val Thr Arg Asp Leu Glu Gly Thr Tyr Leu Cys Arg Ala Arg Ser Thr Gln Gly Glu Val Thr Arg Glu Val Thr Val Asn Val Thr Ser Gly Ser Ser Ala Ser Pro Thr Ser 470 Pro Lys Val Phe Pro Leu Ser Leu Asp Ser Thr Pro Gln Asp Gly Asn Val Val Val Ala Cys Leu Val Gln Gly Phe Phe Pro Gln Glu Pro Leu Ser Val Thr Trp Ser Glu Ser Gly Gln Asn Val Thr Ala Arg Asn Phe 520 Pro Pro Ser Gln Asp Ala Ser Gly Asp Leu Tyr Thr Thr Ser Ser Gln Leu Thr Leu Pro Ala Thr Gln Cys Pro Asp Gly Lys Ser Val Thr Cys 545 His Val Lys His Tyr Thr Asn Ser Ser Gln Asp Val Thr Val Pro Cys 570 Arg Val Pro Pro Pro Pro Cys Cys His Pro Arg Leu Ser Leu His

585

Arg Pro Ala Leu Glu Asp Leu Leu Gly Ser Glu Ala Asn Leu Thr 595 600 605

Cys Thr Leu Thr Gly Leu Arg Asp Ala Ser Gly Ala Thr Phe Thr Trp 610 615 620

Thr Pro Ser Ser Gly Lys Ser Ala Val Gln Gly Pro Pro Glu Arg Asp 625 630 635 640

Leu Cys Gly Cys Tyr Ser Val Ser Arg Val Leu Pro Gly Cys Ala Gln 645 650 655

Pro Trp Asn His Gly Glu Thr Phe Thr Cys Thr Ala Ala His Pro Glu 660 665 . 670

Leu Lys Thr Pro Leu Thr Ala Asn Ile Thr Lys Ser Gly Asn Thr Phe 675 680 685

Arg Pro Glu Val His Leu Leu Pro Pro Pro Ser Glu Glu Leu Ala Leu 690 695 700

Asn Glu Leu Val Thr Leu Thr Cys Leu Ala Arg Gly Phe Ser Pro Lys 705 710 715 720

Asp Val Leu Val Arg Trp Leu Gln Gly Ser Gln Glu Leu Pro Arg Glu 725 730 735

Lys Tyr Leu Thr Trp Ala Ser Arg Gln Glu Pro Ser Gln Gly Thr Thr 740 745 750

Thr Tyr Ala Val Thr Ser Ile Leu Arg Val Ala Ala Glu Asp Trp Lys
755 760 765

Lys Gly Glu Thr Phe Ser Cys Met Val Gly His Glu Ala Leu Pro Leu 770 780

Ala Phe Thr Gln Lys Thr Ile Asp Arg Leu Ala Gly Lys Pro Thr His 785 790 795 800

Ile Asn Val Ser Val Val Met Ala Glu Ala Asp Gly Thr Cys Tyr Arg 805 810 815

Ser Glu Lys Asp Glu Leu 820

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<211> 159

<212> PRT

<213> Homo sapiens

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Asn Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser Ser 35 40 45

Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val 50 60

Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg 65 70 75 80

Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro 85 90 95

Thr Glu Val Glu Leu Asp Asn Gln Ile Val Thr Ala Thr Gln Ser Asn 100 105 110

Ile Cys Asp Glu Asp Ser Ala Thr Glu Thr Cys Tyr Thr Tyr Asp Arg 115 120 125

Asn Lys Cys Tyr Thr Ala Val Val Pro Leu Val Tyr Gly Gly Glu Thr 130 135 140

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<210> 51

<211> 602

<212> PRT

<213> Homo sapiens

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Gly Asn Ser Val Ser Ile Thr Cys Tyr Tyr Pro Pro Thr Ser Val Asn 35 40 45

Arg Thr Arg Lys Tyr Trp Cys Arg Gln Gly Ala Arg Gly Gly Cys Ile

Thr Leu Ile Ser Ser Glu Gly Tyr Val Ser Ser Lys Tyr Ala Gly Arg
65 70 75 80

Ala Asn Leu Thr Asn Phe Pro Glu Asn Gly Thr Phe Val Val Asn Ile 85 90 95

Ala Gln Leu Ser Gln Asp Asp Ser Gly Arg Tyr Lys Cys Gly Leu Gly
100 105 110

Ile Asn Ser Arg Gly Leu Ser Phe Asp Val Ser Leu Glu Val Ser Gln 120 Gly Pro Gly Leu Leu Asn Asp Thr Lys Val Tyr Thr Val Asp Leu Gly Arg Thr Val Thr Ile Asn Cys Pro Phe Lys Thr Glu Asn Ala Gln Lys 150 155 Arg Lys Ser Leu Tyr Lys Gln Ile Gly Leu Tyr Pro Val Leu Val Ile Asp Ser Ser Gly Tyr Val Asn Pro Asn Tyr Thr Gly Arg Ile Arg Leu Asp Ile Gln Gly Thr Gly Gln Leu Leu Phe Ser Val Val Ile Asn Gln Leu Arg Leu Ser Asp Ala Gly Gln Tyr Leu Cys Gln Ala Gly Asp Asp 215 Ser Asn Ser Asn Lys Lys Asn Ala Asp Leu Gln Val Leu Lys Pro Glu 230 Pro Glu Leu Val Tyr Glu Asp Leu Arg Gly Ser Val Thr Phe Cys Ala 250 Leu Gly Pro Glu Val Ala Asn Val Ala Lys Phe Leu Cys Arg Gln Ser Ser Gly Glu Asn Cys Asp Val Val Val Asn Thr Leu Gly Lys Arg Ala Pro Ala Phe Glu Gly Arg Ile Leu Leu Asn Pro Gln Asp Lys Asp Gly Ser Phe Ser Val Val Ile Thr Gly Leu Arg Lys Glu Asp Ala Gly Arg Tyr Leu Cys Gly Ala Ser Asp Gly Gln Leu Gln Glu Gly Ser Pro Ile Gln Ala Trp Gln Leu Phe Val Asn Glu Glu Ser Thr Ile Pro Arg Ser Pro Thr Val Val Lys Gly Val Ala Gly Ser Ser Val Ala Val Leu Cys Pro Tyr Asn Arg Lys Glu Ser Lys Ser Ile Lys Tyr Trp Cys Leu Trp

385 390 395 400

Trp Val Lys Ala Gln Tyr Glu Gly Arg Leu Ser Leu Leu Glu Glu Pro
405 410 415

Glu Gly Ala Gln Asn Gly Arg Cys Pro Leu Leu Val Asp Ser Glu Gly

375

Gly Asn Gly Thr Phe Thr Val Ile Leu Asn Gln Leu Thr Ser Arg Asp 420 425 430

Ala Gly Phe Tyr Trp Cys Leu Thr Asn Gly Asp Thr Leu Trp Arg Thr 435 440 445

Thr Val Glu Ile Lys Ile Ile Glu Gly Glu Pro Asn Leu Lys Val Pro 450 455 460

Gly Asn Val Thr Ala Val Leu Gly Glu Thr Leu Lys Val Pro Cys Phe 465 470 475 480

Pro Cys Lys Phe Ser Ser Tyr Glu Lys Tyr Trp Cys Lys Trp Asn Asn 485 490 495

Thr Gly Cys Gln Ala Leu Pro Ser Gln Asp Glu Gly Pro Ser Lys Ala
500 505 510

Phe Val Asn Cys Asp Glu Asn Ser Arg Leu Val Ser Leu Thr Leu Asn 515 520 525

Leu Val Thr Arg Ala Asp Glu Gly Trp Tyr Trp Cys Gly Val Lys Gln 530 535 540

Gly Phe Tyr Gly Glu Thr Ala Ala Val Tyr Val Ala Val Glu Glu Arg 545 550 555 560

Lys Ala Ala Gly Ser Arg Asp Val Ser Leu Ala Lys Ala Asp Ala Ala 565 570 575

Pro Asp Glu Lys Val Leu Asp Ser Gly Phe Arg Glu Ile Glu Asn Lys
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Ala Ile Gln Asp Pro Arg Leu Phe Ala Glu 595 600

<210> 52

<211> 2533

<212> DNA

<213> Homo sapiens

<400> 52

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accatcacgt gcctggtgac gggcttctct cccgcggacg tcttcgtgca gtggatgcag 1560
agggggcagc ccttgtcccc ggagaagtat gtgaccagcg ccccaatgcc tgagccccag 1620
gccccaggcc ggtacttcgc ccacagcatc ctgaccgtgt ccgaagagga atggaacacg 1680
ggggagacet acacetgegt ggtggeeeat gaggeeetge eeaacagggt cacegagagg 1740
acceptedaca adtecaccea gagagagata agcaccacca agagagactt tagagaaccta 1800
tgggccaccq cetecacett categteete tteeteetga geetetteta cagtaccace 1860
                                                                  1884
gtcaccttgt tcaaggtgaa atga
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<210> 62

<211> 454

<212> PRT

<213> Homo sapiens

<400> 62

Gly Ser Ala Ser Ala Pro Thr Leu Phe Pro Leu Val Ser Cys Glu Asn 1 10 15

Ser Pro Ser Asp Thr Ser Ser Val Ala Val Gly Cys Leu Ala Gln Asp

Phe Leu Pro Asp Ser Ile Thr Phe Ser Trp Lys Tyr Lys Asn Asn Ser 35 40 45

Asp Ile Ser Ser Thr Arg Gly Phe Pro Ser Val Leu Arg Gly Gly Lys
50 60

Tyr Ala Ala Thr Ser Gln Val Leu Leu Pro Ser Lys Asp Val Met Gln 65 70 75 80

Gly Thr Asp Glu His Val Val Cys Lys Val Gln His Pro Asn Gly Asn 90 95

Lys Glu Lys Asn Val Pro Leu Pro Val Ile Ala Glu Leu Pro Pro Lys
100 105 110

Val Ser Val Phe Val Pro Pro Arg Asp Gly Phe Phe Gly Asn Pro Arg 115 120 125 Ser Lys Ser Lys Leu Ile Cys Gln Ala Thr Gly Phe Ser Pro Arg Gln 135 Ile Gln Val Ser Trp Leu Arg Glu Gly Lys Gln Val Gly Ser Gly Val Thr Thr Asp Gln Val Gln Ala Glu Ala Lys Glu Ser Gly Pro Thr Thr 165 Tyr Lys Val Thr Ser Thr Leu Thr Ile Lys Glu Ser Asp Trp Leu Ser 185 Gln Ser Met Phe Thr Cys Arg Val Asp His Arg Gly Leu Thr Phe Gln Gln Asn Ala Ser Ser Met Cys Val Pro Asp Gln Asp Thr Ala Ile Arg Val Phe Ala Ile Pro Pro Ser Phe Ala Ser Ile Phe Leu Thr Lys Ser Thr Lys Leu Thr Cys Leu Val Thr Asp Leu Thr Thr Tyr Asp Ser Val Thr Ile Ser Trp Thr Arg Gln Asn Gly Glu Ala Val Lys Thr His Thr 265 260 Asn Ile Ser Glu Ser His Pro Asn Ala Thr Phe Ser Ala Val Gly Glu 280 Ala Ser Ile Cys Glu Asp Asp Trp Asn Ser Gly Glu Arg Phe Thr Cys Thr Val Thr His Thr Asp Leu Pro Ser Pro Leu Lys Gln Thr Ile Ser Arg Pro Lys Gly Val Ala Leu His Arg Pro Asp Val Tyr Leu Leu Pro Pro Ala Arg Glu Gln Leu Asn Leu Arg Glu Ser Ala Thr Ile Thr Cys Leu Val Thr Gly Phe Ser Pro Ala Asp Val Phe Val Gln Trp Met Gln 360 Arg Gly Gln Pro Leu Ser Pro Glu Lys Tyr Val Thr Ser Ala Pro Met Pro Glu Pro Gln Ala Pro Gly Arg Tyr Phe Ala His Ser Ile Leu Thr Val Ser Glu Glu Glu Trp Asn Thr Gly Glu Thr Tyr Thr Cys Val Val 405 Ala His Glu Ala Leu Pro Asn Arg Val Thr Glu Arg Thr Val Asp Lys 425 420

Ser Thr Gly Lys Pro Thr Leu Tyr Asn Val Ser Leu Val Met Ser Asp 435 440 445

Thr Ala Gly Thr Cys Tyr 450

<210> 63

<211> 532

<212> PRT

<213> Homo sapiens

<400> 63

Met Ala Pro Ser Ser Pro Arg Pro Ala Leu Pro Ala Leu Leu Val Leu 1 5 10 15

Leu Gly Ala Leu Phe Pro Gly Pro Gly Asn Ala Gln Thr Ser Val Ser 20 25 30

Pro Ser Lys Val Ile Leu Pro Arg Gly Gly Ser Val Leu Val Thr Cys 35 40 45

Ser Thr Ser Cys Asp Gln Pro Lys Leu Leu Gly Ile Glu Thr Pro Leu 50 55 60

Pro Lys Lys Glu Leu Leu Pro Gly Asn Asn Arg Lys Val Tyr Glu 65 70 75 80

Leu Ser Asn Val Gln Glu Asp Ser Gln Pro Met Cys Tyr Ser Asn Cys 85 90 95

Pro Asp Gly Gln Ser Thr Ala Lys Thr Phe Leu Thr Val Tyr Trp Thr

Pro Glu Arg Val Glu Leu Ala Pro Leu Pro Ser Trp Gln Pro Val Gly 115 120 125

Lys Asn Leu Thr Leu Arg Cys Gln Val Glu Gly Gly Ala Pro Arg Ala 130 135 140

Asn Leu Thr Val Val Leu Leu Arg Gly Glu Lys Glu Leu Lys Arg Glu 145 150 155 160

Pro Ala Val Gly Glu Pro Ala Glu Val Thr Thr Thr Val Leu Val Arg 165 170 175

Arg Asp His His Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu Asp Leu 180 185 190

Arg Pro Gln Gly Leu Glu Leu Phe Glu Asn Thr Ser Ala Pro Tyr Gln
195 200 205

Leu Gln Thr Phe Val Leu Pro Ala Thr Pro Pro Gln Leu Val Ser Pro 210 215 220

Arg Val Leu Glu Val Asp Thr Gln Gly Thr Val Val Cys Ser Leu Asp 225 230 235 240

Gly Leu Phe Pro Val Ser Glu Ala Gln Val His Leu Ala Leu Gly Asp 245 250 255

Gln Arg Leu Asn Pro Thr Val Thr Tyr Gly Asn Asp Ser Phe Ser Ala
260 265 270

Lys Ala Ser Val Ser Val Thr Ala Glu Asp Glu Gly Thr Gln Arg Leu 275 280 285

Thr Cys Ala Val Ile Leu Gly Asn Gln Ser Gln Glu Thr Leu Gln Thr 290 295 300

Val Thr Ile Tyr Ser Phe Pro Ala Pro Asn Val Ile Leu Thr Lys Pro 305 310 315 320

Glu Val Ser Glu Gly Thr Glu Val Thr Val Lys Cys Glu Ala His Pro 325 330 335

Arg Ala Lys Val Thr Leu Asn Gly Val Pro Ala Gln Pro Leu Gly Pro 340 345 350

Arg Ala Gln Leu Leu Lys Ala Thr Pro Glu Asp Asn Gly Arg Ser 355 360 365

Phe Ser Cys Ser Ala Thr Leu Glu Val Ala Gly Gln Leu Ile His Lys 370 375 380

Asn Gln Thr Arg Glu Leu Arg Val Leu Tyr Gly Pro Arg Leu Asp Glu 385 390 395 400

Arg Asp Cys Pro Gly Asn Trp Thr Trp Pro Glu Asn Ser Gln Gln Thr 405 410 415

Pro Met Cys Gln Ala Trp Gly Asn Pro Leu Pro Glu Leu Lys Cys Leu 420 425 430

Lys Asp Gly Thr Phe Pro Leu Pro Ile Gly Glu Ser Val Thr Val Thr 435 440 445

Arg Asp Leu Glu Gly Thr Tyr Leu Cys Arg Ala Arg Ser Thr Gln Gly 450 455 460

Glu Val Thr Arg Glu Val Thr Val Asn Val Leu Ser Pro Arg Tyr Glu 465 470 475 480

Ile Val Ile Ile Thr Val Val Ala Ala Val Ile Met Gly Thr Ala
485 490 495

Gly Leu Ser Thr Tyr Leu Tyr Asn Arg Gln Arg Lys Ile Lys Lys Tyr 500 505 510

Arg Leu Gln Gln Ala Gln Lys Gly Thr Pro Met Lys Pro Asn Thr Gln 515 520 525

Ala Thr Pro Pro 530 <210> 64

<211> 275

<212> PRT

<213> Homo sapiens

<400> 64

Met Ser Ser Phe Gly Tyr Arg Thr Leu Thr Val Ala Leu Phe Thr Leu

1 5 10 15

Ile Cys Cys Pro Gly Ser Asp Glu Lys Val Phe Glu Val His Val Arg 20 25 30

Pro Lys Lys Leu Ala Val Glu Pro Lys Gly Ser Leu Glu Val Asn Cys 35 40 45

Ser Thr Thr Cys Asn Gln Pro Glu Val Gly Gly Leu Glu Thr Ser Leu 50 55 60

Asp Lys Ile Leu Leu Asp Glu Gln Ala Gln Trp Lys His Tyr Leu Val 65 70 75 80

Ser Asn Ile Ser His Asp Thr Val Leu Gln Cys His Phe Thr Cys Ser 85 90 95

Gly Lys Gln Glu Ser Met Asn Ser Asn Val Ser Val Tyr Gln Pro Pro 100 105 110

Arg Gln Val Ile Leu Thr Leu Gln Pro Thr Leu Val Ala Val Gly Lys
115 120 125

Ser Phe Thr Ile Glu Cys Arg Val Pro Thr Val Glu Pro Leu Asp Ser 130 135 140

Leu Thr Leu Phe Leu Phe Arg Gly Asn Glu Thr Leu His Tyr Glu Thr 145 150 155 160

Phe Gly Lys Ala Ala Pro Ala Pro Gln Glu Ala Thr Ala Thr Phe Asn 165 170 175

Ser Thr Ala Asp Arg Glu Asp Gly His Arg Asn Phe Ser Cys Leu Ala 180 185 190

Val Leu Asp Leu Met Ser Arg Gly Gly Asn Ile Phe His Lys His Ser 195 200 205

Ala Pro Lys Met Leu Glu Ile Tyr Glu Pro Val Ser Asp Ser Gln Met 210 215 220

Val Ile Ile Val Thr Val Val Ser Val Leu Leu Ser Leu Phe Val Thr 225 230 235 240

Ser Val Leu Leu Cys Phe Ile Phe Gly Gln His Leu Arg Gln Gln Arg 245 250 255

Met Gly Thr Tyr Gly Val Arg Ala Ala Trp Arg Arg Leu Pro Gln Ala 260 265 270

Phe Arg Pro 275

<210> 65

<211> 547

<212> PRT

<213> Homo sapiens

<400> 65

Met Ala Thr Met Val Pro Ser Val Leu Trp Pro Arg Ala Cys Trp Thr 1 5 10 15

Leu Leu Val Cys Cys Leu Leu Thr Pro Gly Val Gln Gly Gln Glu Phe 20 25 30

Leu Leu Arg Val Glu Pro Gln Asn Pro Val Leu Ser Ala Gly Gly Ser
35 40 45

Leu Phe Val Asn Cys Ser Thr Asp Cys Pro Ser Ser Glu Lys Ile Ala 50 55 60

Leu Glu Thr Ser Leu Ser Lys Glu Leu Val Ala Ser Gly Met Gly Trp
65 70 75 80

Ala Ala Phe Asn Leu Ser Asn Val Thr Gly Asn Ser Arg Ile Leu Cys
85 90 95

Ser Val Tyr Cys Asn Gly Ser Gln Ile Thr Gly Ser Ser Asn Ile Thr 100 105 110

Val Tyr Gly Leu Pro Glu Arg Val Glu Leu Ala Pro Leu Pro Pro Trp 115 120 125

Gln Pro Val Gly Gln Asn Phe Thr Leu Arg Cys Gln Val Glu Gly Gly 130 135 140

Ser Pro Arg Thr Ser Leu Thr Val Val Leu Leu Arg Trp Glu Glu Glu 145 150 155 160

Leu Ser Arg Gln Pro Ala Val Glu Glu Pro Ala Glu Val Thr Ala Thr
165 170 175

Val Leu Ala Ser Arg Asp Asp His Gly Ala Pro Phe Ser Cys Arg Thr 180 185 190

Glu Leu Asp Met Gln Pro Gln Gly Leu Gly Leu Phe Val Asn Thr Ser 195 200 205

Ala Pro Arg Gln Leu Arg Thr Phe Val Leu Pro Val Thr Pro Pro Arg 210 215 220

Leu Val Ala Pro Arg Phe Leu Glu Val Glu Thr Ser Trp Pro Val Asp 225 230 235 240

Cys Thr Leu Asp Gly Leu Phe Pro Ala Ser Glu Ala Gln Val Tyr Leu 245 250 255

Ala Leu Gly Asp Gln Met Leu Asn Ala Thr Val Met Asn His Gly Asp

Thr Leu Thr Ala Thr Ala Thr Ala Thr Ala Arg Ala Asp Gln Glu Gly 280

Ala Arg Glu Ile Val Cys Asn Val Thr Leu Gly Gly Glu Arg Arg Glu

Ala Arg Glu Asn Leu Thr Val Phe Ser Phe Leu Gly Pro Ile Val Asn 315

Leu Ser Glu Pro Thr Ala His Glu Gly Ser Thr Val Thr Val Ser Cys

Met Ala Gly Ala Arg Val Gln Val Thr Leu Asp Gly Val Pro Ala Ala 345

Ala Pro Gly Gln Pro Ala Gln Leu Gln Leu Asn Ala Thr Glu Ser Asp 360 355

Asp Gly Arg Ser Phe Phe Cys Ser Ala Thr Leu Glu Val Asp Gly Glu 375

Phe Leu His Arq Asn Ser Ser Val Gln Leu Arg Val Leu Tyr Gly Pro 390 395

Lys Ile Asp Arq Ala Thr Cys Pro Gln His Leu Lys Trp Lys Asp Lys

Thr Arg His Val Leu Gln Cys Gln Ala Arg Gly Asn Pro Tyr Pro Glu

Leu Arg Cys Leu Lys Glu Gly Ser Ser Arg Glu Val Pro Val Gly Ile

Pro Phe Phe Val Asn Val Thr His Asn Gly Thr Tyr Gln Cys Gln Ala 455

Ser Ser Ser Arg Gly Lys Tyr Thr Leu Val Val Wet Asp Ile Glu

Ala Gly Ser Ser His Phe Val Pro Val Phe Val Ala Val Leu Leu Thr

Leu Gly Val Val Thr Ile Val Leu Ala Leu Met Tyr Val Phe Arg Glu 500

His Gln Arg Ser Gly Ser Tyr His Val Arg Glu Glu Ser Thr Tyr Leu 520

Pro Leu Thr Ser Met Gln Pro Thr Glu Ala Met Gly Glu Glu Pro Ser 530 535

Arg Ala Glu

<210> 66

<211> 577

<212> PRT

<213> Homo sapiens

<400> 66

Gly Val Pro Glu Glu Leu Phe Glu Val Ser Ile Trp Pro Ser Gln Ala 1 5 10 15

Leu Val Glu Phe Gly Gln Ser Leu Val Val Asn Cys Ser Thr Thr Cys 20 25 30

Pro Asp Pro Gly Pro Ser Gly Ile Glu Thr Phe Leu Lys Lys Thr Gln 35 40 45

Val Gly Lys Gly Pro Gln Trp Lys Glu Phe Leu Leu Glu Asp Val Thr 50 55 60

Glu Asn Ser Ile Leu Gln Cys Phe Phe Ser Cys Ala Gly Ile Gln Lys 65 70 75 80

Asp Thr Ser Leu Gly Ile Thr Val Tyr Gln Pro Pro Glu Gln Val Ile 85 90 95

Leu Glu Leu Gln Pro Ala Trp Val Ala Val Asp Glu Ala Phe Thr Val 100 105 110

Lys Cys His Val Pro Ser Val Ala Pro Leu Glu Ser Leu Thr Leu Ala 115 120 125

Leu Leu Gln Gly Asn Gln Glu Leu His Arg Lys Asn Phe Thr Ser Leu 130 135 140

Ala Val Ala Ser Gln Arg Ala Glu Val Ile Ile Ser Val Arg Ala Gln 145 150 155 160

Lys Glu Asn Asp Arg Cys Asn Ser Ser Cys His Ala Glu Leu Asp Leu 165 170 175

Ser Leu Gln Gly Gly Arg Leu Phe Gln Gly Ser Ser Pro Ile Arg Ile 180 185 190

Val Arg Ile Phe Glu Phe Ser Gln Ser Pro His Ile Trp Val Ser Ser 195 200 205

Leu Leu Glu Ala Gly Met Ala Glu Thr Val Ser Cys Glu Val Ala Arg 210 215 220

Val Phe Pro Ala Lys Glu Val Met Phe His Met Phe Leu Glu Asp Gln 225 230 235 240

Glu Leu Ser Ser Phe Leu Ser Trp Glu Gly Asp Thr Ala Trp Ala Asn 245 250 255

Ala Thr Ile Arg Thr Met Glu Ala Gly Asp Gln Glu Leu Ser Cys Phe 260 265 270 Ala Ser Leu Gly Ala Met Glu Gln Lys Thr Arg Lys Leu Val His Ser Tyr Asn Lys Trp Pro Gly Ser Ser Phe Phe Ile Arg Val Leu Cys Cys 295 Lys His Arg Val Thr Gly Trp Phe Gly Cys Arg His Pro Cys Cys Pro Leu Leu Gly Met Leu Ser Ser Glu His Glu Ser Ser Ser Phe Ser Gly 330 Phe Pro Pro Pro Ile Leu Glu Leu Lys Glu Ser Tyr Pro Leu Ala Gly Thr Asp Ile Asn Val Thr Cys Ser Gly His Val Leu Thr Ser Pro Ser Pro Thr Leu Arg Leu Gln Gly Ala Pro Asp Leu Pro Ala Gly Glu Pro 375 Ala Trp Leu Leu Thr Ala Arg Glu Glu Asp Asp Gly Asn Phe Ser 390 Cys Glu Ala Ser Leu Val Val Gln Gly Gln Arg Leu Met Lys Thr Thr 405 Val Ile Gln Leu His Ile Leu Cys Lys Pro Gln Leu Glu Glu Ser Ser Cys Pro Gly Lys Gln Thr Trp Leu Glu Gly Met Glu His Thr Leu Ala Cys Val Pro Lys Gly Asn Pro Ala Pro Ala Leu Val Cys Thr Trp Asn Gly Val Val Phe Asp Leu Glu Val Pro Gln Lys Ala Thr Asn His Thr 470 475 Gly Thr Tyr Arg Tyr Thr Ala Thr Asn Gln Leu Gly Ser Val Ser Lys Asp Ile Ala Val Ile Val Gln Gly Leu Asp Glu Gly Ile Ser Ser Thr Leu Phe Val Ile Ile Thr Val Ala Leu Gly Val Gly Val Ile Thr Ile 515 Ala Leu Tyr Leu Ser Tyr Arg Pro Cys Lys Val Asp Arg Arg Lys Leu 535 Leu Tyr Arg Gln Lys Glu Glu Asp Lys Glu Glu Glu Ser Gln Phe Ala Val Gln Glu Glu Lys Ser Thr Thr His Ile Ile Asp Ser Tyr Leu Ile

<210> 67

<211> 924

<212> PRT

<213> Homo sapiens

<400> 67

Met Pro Gly Pro Ser Pro Gly Leu Arg Arg Ala Leu Leu Gly Leu Trp
1 5 10 15

Ala Ala Leu Gly Leu Gly Leu Phe Gly Leu Ser Ala Val Ser Gln Glu 20 25 30

Pro Phe Trp Ala Asp Leu Gln Pro Arg Val Ala Phe Val Glu Arg Gly 35 40 45

Gly Ser Leu Trp Leu Asn Cys Ser Thr Asn Cys Pro Arg Pro Glu Arg 50 55 60

Gly Gly Leu Glu Thr Ser Leu Arg Arg Asn Gly Thr Gln Arg Gly Leu 65 70 75 80

Arg Trp Leu Ala Arg Gln Leu Val Asp Ile Arg Glu Pro Glu Thr Gln
85 90 95

Pro Val Cys Phe Phe Arg Cys Ala Arg Arg Thr Leu Gln Ala Arg Gly
100 105 110

Leu Ile Arg Thr Phe Gln Arg Pro Asp Arg Val Glu Leu Met Pro Leu 115 120 125

Pro Pro Trp Gln Pro Val Gly Glu Asn Phe Thr Leu Ser Cys Arg Val 130 135 140

Pro Gly Ala Gly Pro Arg Ala Ser Leu Thr Leu Thr Leu Leu Arg Gly 145 150 155 160

Ala Gln Glu Leu Ile Arg Arg Ser Phe Ala Gly Glu Pro Pro Arg Ala 165 170 175

Arg Gly Ala Val Leu Thr Ala Thr Val Leu Ala Arg Arg Glu Asp His 180 185 190

Gly Ala Asn Phe Ser Cys Arg Ala Glu Leu Asp Leu Arg Pro His Gly
195 200 205

Leu Gly Leu Phe Glu Asn Ser Ser Ala Pro Arg Glu Leu Arg Thr Phe 210 215 220

Ser Leu Ser Pro Asp Ala Pro Arg Leu Ala Ala Pro Arg Leu Glu 225 230 235 240

Val Gly Ser Glu Arg Pro Val Ser Cys Thr Leu Asp Gly Leu Phe Pro 245 250 255

Ala Ser Glu Ala Arg Val Tyr Leu Ala Leu Gly Asp Gln Asn Leu Ser 260 265 270 Pro Asp Val Thr Leu Glu Gly Asp Ala Phe Val Ala Thr Ala 275 280 285

Thr Ala Ser Ala Glu Gln Glu Gly Ala Arg Gln Leu Ile Cys Asn Val 290 295 300

Thr Leu Gly Gly Glu Asn Arg Glu Thr Arg Glu Asn Val Thr Ile Tyr 305 310 315 320

Ser Phe Pro Ala Pro Leu Leu Thr Leu Ser Glu Pro Ser Val Ser Glu 325 330 335

Gly Gln Met Val Thr Val Thr Cys Ala Ala Gly Thr Gln Ala Leu Val 340 345 350

Thr Leu Glu Gly Val Pro Ala Ala Val Pro Gly Gln Pro Ala Gln Leu 355 360 365

Gln Leu Asn Ala Thr Glu Asn Asp Asp Arg Arg Ser Phe Phe Cys Asp 370 375 380

Ala Thr Leu Asp Val Asp Gly Glu Thr Leu Ile Lys Asn Arg Ser Ala 385 390 395 400

Glu Leu Arg Val Leu Tyr Ala Pro Arg Leu Asp Asp Ser Asp Cys Pro 405 410 415

Arg Ser Trp Thr Trp Pro Glu Gly Pro Glu Gln Thr Leu Arg Cys Glu
420 425 430

Ala Arg Gly Asn Pro Glu Pro Ser Val His Cys Ala Arg Ser Asp Gly
435 440 445

Gly Ala Val Leu Ala Leu Gly Leu Leu Gly Pro Val Thr Arg Ala Leu 450 455 460

Ser Gly Thr Tyr Arg Cys Lys Ala Ala Asn Asp Gln Gly Glu Ala Val 465 470 475 480

Lys Asp Val Thr Leu Thr Val Glu Tyr Ala Pro Ala Leu Asp Ser Val 485 490 495

Gly Cys Pro Glu Arg Ile Thr Trp Leu Glu Gly Thr Glu Ala Ser Leu 500 505 510

Ser Cys Val Ala His Gly Val Pro Pro Pro Asp Val Ile Cys Val Arg 515 520 525

Ser Gly Glu Leu Gly Ala Val Ile Glu Gly Leu Leu Arg Val Ala Arg 530 535 540

Glu His Ala Gly Thr Tyr Arg Cys Glu Ala Thr Asn Pro Arg Gly Ser 555 560

Ala Ala Lys Asn Val Ala Val Thr Val Glu Tyr Gly Pro Arg Phe Glu 565 570 575 Glu Pro Ser Cys Pro Ser Asn Trp Thr Trp Val Glu Gly Ser Gly Arg 580 585 590

Leu Phe Ser Cys Glu Val Asp Gly Lys Pro Gln Pro Ser Val Lys Cys 595 600 605

Val Gly Ser Gly Gly Ala Thr Glu Gly Val Leu Leu Pro Leu Ala Pro 610 615 620

Pro Asp Pro Ser Pro Arg Ala Pro Arg Ile Pro Arg Val Leu Ala Pro 625 630 635 640

Gly Ile Tyr Val Cys Asn Ala Thr Asn Arg His Gly Ser Val Ala Lys 645 650 655

Thr Val Val Ser Ala Glu Ser Pro Pro Glu Met Asp Glu Ser Thr 660 665 670

Cys Pro Ser His Gln Thr Trp Leu Glu Gly Ala Glu Ala Ser Ala Leu 675 680 685

Ala Cys Ala Ala Arg Gly Arg Pro Ser Pro Gly Val Arg Cys Ser Arg 690 695 700

Glu Gly Ile Pro Trp Pro Glu Gln Gln Arg Val Ser Arg Glu Asp Ala 705 710 715 720

Gly Thr Tyr His Cys Val Ala Thr Asn Ala His Gly Thr Asp Ser Arg
725 730 735

Thr Val Thr Val Gly Val Glu Tyr Arg Pro Val Val Ala Glu Leu Ala 740 745 750

Ala Ser Pro Pro Gly Gly Val Arg Pro Gly Gly Asn Phe Thr Leu Thr 755 760 765

Cys Arg Ala Glu Ala Trp Pro Pro Ala Gln Ile Ser Trp Arg Ala Pro 770 775 780

Pro Gly Ala Leu Asn Ile Gly Leu Ser Ser Asn Asn Ser Thr Leu Ser 785 790 795 800

Val Ala Gly Ala Met Gly Ser His Gly Glu Tyr Glu Cys Ala Arg 805 810 815

Thr Asn Ala His Gly Arg His Ala Arg Arg Ile Thr Val Arg Val Ala 820 825 830

Gly Pro Trp Leu Trp Val Ala Val Gly Gly Ala Ala Gly Gly Ala Ala 835 840 845

Leu Leu Ala Ala Gly Ala Gly Leu Ala Phe Tyr Val Gln Ser Thr Ala 850 855 860

Cys Lys Lys Gly Glu Tyr Asn Val Gln Glu Ala Glu Ser Ser Gly Glu 865 870 875 880 Ala Val Cys Leu Asn Gly Ala Gly Gly Gly Ala Gly Gly Ala Ala Gly 885 890 895

Ala Glu Gly Gly Pro Glu Ala Ala Gly Gly Ala Ala Glu Ser Pro Ala 900 905 910

Glu Gly Glu Val Phe Ala Ile Gln Leu Thr Ser Ala 915 920

<210> 68

<211> 406

<212> PRT

<213> Homo sapiens

<400> 68

Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu 1 5 10 15

Leu Gly Gln Ser Leu Gln Val Lys Pro Leu Gln Val Glu Pro Pro Glu 20 25 30

Pro Val Val Ala Val Ala Leu Gly Ala Ser Arg Gln Leu Thr Cys Arg 35 40 45

Leu Ala Cys Ala Asp Arg Gly Ala Ser Val Gln Trp Arg Gly Leu Asp 50 55 60

Thr Ser Leu Gly Ala Val Gln Ser Asp Thr Gly Arg Ser Val Leu Thr 65 70 75 80

Val Arg Asn Ala Ser Leu Ser Ala Ala Gly Thr Arg Val Cys Val Gly 85 90 95

Ser Cys Gly Gly Arg Thr Phe Gln His Thr Val Gln Leu Leu Val Tyr 100 105 110

Ala Phe Pro Asp Gln Leu Thr Val Ser Pro Ala Ala Leu Val Pro Gly
115 120 125

Asp Pro Glu Val Ala Cys Thr Ala His Lys Val Thr Pro Val Asp Pro 130 135 140

Asn Ala Leu Ser Phe Ser Leu Leu Val Gly Gly Gln Glu Leu Glu Gly 145 150 155 160

Ala Gln Ala Leu Gly Pro Glu Val Gln Glu Glu Glu Glu Glu Pro Gln
165 170 175

Gly Asp Glu Asp Val Leu Phe Arg Val Thr Glu Arg Trp Arg Leu Pro 180 185 190

Pro Leu Gly Thr Pro Val Pro Pro Ala Leu Tyr Cys Gln Ala Thr Met 195 200 205

Arg Leu Pro Gly Leu Glu Leu Ser His Arg Gln Ala Ile Pro Val Leu 210 215 220

His Ser Pro Thr Ser Pro Glu Pro Pro Asp Thr Thr Ser Pro Glu Pro 230 Pro Asn Thr Thr Ser Pro Glu Ser Pro Asp Thr Thr Ser Pro Glu Ser 250 245 Pro Asp Thr Thr Ser Gln Glu Pro Pro Asp Thr Thr Ser Gln Glu Pro 265 Pro Asp Thr Thr Ser Gln Glu Pro Pro Asp Thr Thr Ser Pro Glu Pro Pro Asp Lys Thr Ser Pro Glu Pro Ala Pro Gln Gln Gly Ser Thr His Thr Pro Arg Ser Pro Gly Ser Thr Arg Thr Arg Arg Pro Glu Ile Ser Gln Ala Gly Pro Thr Gln Gly Glu Val Ile Pro Thr Gly Ser Ser Lys 330 325 Pro Ala Gly Asp Gln Leu Pro Ala Ala Leu Trp Thr Ser Ser Ala Val Leu Gly Leu Leu Leu Leu Ala Leu Pro Thr Tyr His Leu Trp Lys Arg 360 Cys Arg His Leu Ala Glu Asp Asp Thr His Pro Pro Ala Ser Leu Arg Leu Leu Pro Gln Val Ser Ala Trp Ala Gly Leu Arg Gly Thr Gly Gln Val Gly Ile Ser Pro Ser 405 <210> 69 <211> 739 <212> PRT <213> Homo sapiens <400> 69 Met Pro Gly Lys Met Val Val Ile Leu Gly Ala Ser Asn Ile Leu Trp Ile Met Phe Ala Ala Ser Gln Ala Phe Lys Ile Glu Thr Thr Pro Glu 25 Ser Arg Tyr Leu Ala Gln Ile Gly Asp Ser Val Ser Leu Thr Cys Ser Thr Thr Gly Cys Glu Ser Pro Phe Phe Ser Trp Arg Thr Gln Ile Asp

Ser Pro Leu Asn Gly Lys Val Thr Asn Glu Gly Thr Thr Ser Thr Leu

Thr Met Asn Pro Val Ser Phe Gly Asn Glu His Ser Tyr Leu Cys Thr 85 90 95

Ala Thr Cys Glu Ser Arg Lys Leu Glu Lys Gly Ile Gln Val Glu Ile 100 105 110

Tyr Ser Phe Pro Lys Asp Pro Glu Ile His Leu Ser Gly Pro Leu Glu 115 120 125

Ala Gly Lys Pro Ile Thr Val Lys Cys Ser Val Ala Asp Val Tyr Pro 130 135 140

Phe Asp Arg Leu Glu Ile Asp Leu Leu Lys Gly Asp His Leu Met Lys 145 150 155 160

Ser Gln Glu Phe Leu Glu Asp Ala Asp Arg Lys Ser Leu Glu Thr Lys 165 170 175

Ser Leu Glu Val Thr Phe Thr Pro Val Ile Glu Asp Ile Gly Lys Val

Leu Val Cys Arg Ala Lys Leu His Ile Asp Glu Met Asp Ser Val Pro 195 200 205

Thr Val Arg Gln Ala Val Lys Glu Leu Gln Val Tyr Ile Ser Pro Lys 210 215 220

Asn Thr Val Ile Ser Val Asn Pro Ser Thr Lys Leu Gln Glu Gly Gly 225 230 235 240

Ser Val Thr Met Thr Cys Ser Ser Glu Gly Leu Pro Ala Pro Glu Ile 245 250 255

Phe Trp Ser Lys Lys Leu Asp Asn Gly Asn Leu Gln His Leu Ser Gly 260 265 270

Asn Ala Thr Leu Thr Leu Ile Ala Met Arg Met Glu Asp Ser Gly Ile 275 280 285

Tyr Val Cys Glu Gly Val Asn Leu Ile Gly Lys Asn Arg Lys Glu Val 290 295 300

Glu Leu Ile Val Gln Glu Lys Pro Phe Thr Val Glu Ile Ser Pro Gly 305 310 315 320

Pro Arg Ile Ala Ala Gln Ile Gly Asp Ser Val Met Leu Thr Cys Ser 325 330 335

Val Met Gly Cys Glu Ser Pro Ser Phe Ser Trp Arg Thr Gln Ile Asp 340 345 350

Ser Pro Leu Ser Gly Lys Val Arg Ser Glu Gly Thr Asn Ser Thr Leu 355 360 365

Thr Leu Ser Pro Val Ser Phe Glu Asn Glu His Ser Tyr Leu Cys Thr 370 375 380

Val Thr Cys Gly His Lys Lys Leu Glu Lys Gly Ile Gln Val Glu Leu Tyr Ser Phe Pro Arg Asp Pro Glu Ile Glu Met Ser Gly Gly Leu Val Asn Gly Ser Ser Val Thr Val Ser Cys Lys Val Pro Ser Val Tyr Pro Leu Asp Arg Leu Glu Ile Glu Leu Leu Lys Gly Glu Thr Ile Leu Glu Asn Ile Glu Phe Leu Glu Asp Thr Asp Met Lys Ser Leu Glu Asn Lys 455 Ser Leu Glu Met Thr Phe Ile Pro Thr Ile Glu Asp Thr Gly Lys Ala Leu Val Cys Gln Ala Lys Leu His Ile Asp Asp Met Glu Phe Glu Pro Lys Gln Arg Gln Ser Thr Gln Thr Leu Tyr Val Asn Val Ala Pro Arg Asp Thr Thr Val Leu Val Ser Pro Ser Ser Ile Leu Glu Glu Gly Ser 515 Ser Val Asn Met Thr Cys Leu Ser Gln Gly Phe Pro Ala Pro Lys Ile 535 Leu Trp Ser Arg Gln Leu Pro Asn Gly Glu Leu Gln Pro Leu Ser Glu Asn Ala Thr Leu Thr Leu Ile Ser Thr Lys Met Glu Asp Ser Gly Val Tyr Leu Cys Glu Gly Ile Asn Gln Ala Gly Arg Ser Arg Lys Glu Val Glu Leu Ile Ile Gln Val Thr Pro Lys Asp Ile Lys Leu Thr Ala Phe Pro Ser Glu Ser Val Lys Glu Gly Asp Thr Val Ile Ile Ser Cys Thr 615 Cys Gly Asn Val Pro Glu Thr Trp Ile Ile Leu Lys Lys Lys Ala Glu Thr Gly Asp Thr Val Leu Lys Ser Ile Asp Gly Ala Tyr Thr Ile Arg Lys Ala Gln Leu Lys Asp Ala Gly Val Tyr Glu Cys Glu Ser Lys Asn Lys Val Gly Ser Gln Leu Arg Ser Leu Thr Leu Asp Val Gln Gly Arg

680

Glu Asn Asn Lys Asp Tyr Phe Ser Pro Glu Leu Leu Val Leu Tyr Phe 690 695 700

Ala Ser Ser Leu Ile Ile Pro Ala Ile Gly Met Ile Ile Tyr Phe Ala 705 710 715 720

Arg Lys Ala Asn Met Lys Gly Ser Tyr Ser Leu Val Glu Ala Gln Lys 725 730 735

Ser Lys Val

<210> 70

<211> 537

<212> PRT

<213> Mus musculus

<400> 70

Met Ala Ser Thr Arg Ala Lys Pro Thr Leu Pro Leu Leu Leu Ala Leu 1 5 10 15

Val Thr Val Val Ile Pro Gly Pro Gly Asp Ala Gln Val Ser Ile His
20 25 30

Pro Arg Glu Ala Phe Leu Pro Gln Gly Gly Ser Val Gln Val Asn Cys 35 40 45

Ser Ser Ser Cys Lys Glu Asp Leu Ser Leu Gly Leu Glu Thr Gln Trp 50 55 60

Leu Lys Asp Glu Leu Glu Ser Gly Pro Asn Trp Lys Leu Phe Glu Leu 65 70 75 80

Ser Glu Ile Gly Glu Asp Ser Ser Pro Leu Cys Phe Glu Asn Cys Gly 85 90 95

Thr Val Gln Ser Ser Ala Ser Ala Thr Ile Thr Val Tyr Ser Phe Pro 100 105 110

Glu Ser Val Glu Leu Arg Pro Leu Pro Ala Trp Gln Gln Val Gly Lys 115 120 125

Asp Leu Thr Leu Arg Cys His Val Asp Gly Gly Ala Pro Arg Thr Gln 130 135 140

Leu Ser Ala Val Leu Leu Arg Gly Glu Glu Ile Leu Ser Arg Gln Pro 145 150 155 160

Val Gly Gly His Pro Lys Asp Pro Lys Glu Ile Thr Phe Thr Val Leu 165 170 175

Ala Ser Arg Gly Asp His Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu 180 185 190

Asp Leu Arg Pro Gln Gly Leu Ala Leu Phe Ser Asn Val Ser Glu Ala 195 200 205 Arg Ser Leu Arg Thr Phe Asp Leu Pro Ala Thr Ile Pro Lys Leu Asp 215 Thr Pro Asp Leu Leu Glu Val Gly Thr Gln Gln Lys Leu Phe Cys Ser 230 235 Leu Glu Gly Leu Phe Pro Ala Ser Glu Ala Arg Ile Tyr Leu Glu Leu 250 Gly Gly Gln Met Pro Thr Gln Glu Ser Thr Asn Ser Ser Asp Ser Val Ser Ala Thr Ala Leu Val Glu Val Thr Glu Glu Phe Asp Arg Thr Leu 275 Pro Leu Arg Cys Val Leu Glu Leu Ala Asp Gln Ile Leu Glu Thr Gln 295 Arg Thr Leu Thr Val Tyr Asn Phe Ser Ala Pro Val Leu Thr Leu Ser Gln Leu Glu Val Ser Glu Gly Ser Gln Val Thr Val Lys Cys Glu Ala His Ser Gly Ser Lys Val Val Leu Leu Ser Gly Val Glu Pro Arg Pro Pro Thr Pro Gln Val Gln Phe Thr Leu Asn Ala Ser Ser Glu Asp His 360 Lys Arg Ser Phe Phe Cys Ser Ala Ala Leu Glu Val Ala Gly Lys Phe 375 370 Leu Phe Lys Asn Gln Thr Leu Glu Leu His Val Leu Tyr Gly Pro Arg 390 Leu Asp Glu Thr Asp Cys Leu Gly Asn Trp Thr Trp Gln Glu Gly Ser 405 Gln Gln Thr Leu Lys Cys Gln Ala Trp Gly Asn Pro Ser Pro Lys Met Thr Cys Arg Arg Lys Ala Asp Gly Ala Leu Leu Pro Ile Gly Val Val Lys Ser Val Lys Gln Glu Met Asn Gly Thr Tyr Val Cys His Ala Phe 455 Ser Ser His Gly Asn Val Thr Arg Asn Val Tyr Leu Thr Val Leu Tyr 470 475 His Ser Gln Asn Asn Trp Thr Ile Ile Leu Val Pro Val Leu Leu Val Ile Val Gly Leu Val Met Ala Ala Ser Tyr Val Tyr Asn Arg Gln 505

Arg Lys Ile Arg Ile Tyr Lys Leu Gln Lys Ala Gln Glu Glu Ala Ile 515 520 525

Lys Leu Lys Gly Gln Ala Pro Pro Pro 530 535

<210> 71

<211> 537

<212> PRT

<213> Mus musculus

<400> 71

Met Ala Ser Thr Arg Ala Lys Pro Thr Leu Pro Leu Leu Leu Ala Leu
1 5 10 15

Val Thr Val Val Ile Pro Gly Pro Gly Asp Ala Gln Val Ser Ile His
20 25 30

Pro Arg Glu Ala Phe Leu Pro Gln Gly Gly Ser Val Gln Val Asn Cys
35 40 45

Ser Ser Ser Cys Lys Glu Asp Leu Ser Leu Gly Leu Glu Thr Gln Trp 50 60

Leu Lys Asp Glu Leu Glu Ser Gly Pro Asn Trp Lys Leu Phe Glu Leu 65 70 75 80

Ser Glu Ile Gly Glu Asp Ser Ser Pro Leu Cys Phe Glu Asn Cys Gly 85 90 95

Thr Val Gln Ser Ser Ala Ser Ala Thr Ile Thr Val Tyr Ser Phe Pro 100 105 110

Glu Ser Val Glu Leu Arg Pro Leu Pro Ala Trp Gln Gln Val Gly Lys 115 120 125

Asp Leu Thr Leu Arg Cys His Val Asp Gly Gly Ala Pro Arg Thr Gln 130 135 140

Leu Ser Ala Val Leu Leu Arg Gly Glu Glu Ile Leu Ser Arg Gln Pro 145 150 155 160

Val Gly Gly His Pro Lys Asp Pro Lys Glu Ile Thr Phe Thr Val Leu 165 170 175

Ala Ser Arg Gly Asp His Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu 180 185 190

Asp Leu Arg Pro Gln Gly Leu Ala Leu Phe Ser Asn Val Ser Glu Ala 195 200 205

Arg Ser Leu Arg Thr Phe Asp Leu Pro Ala Thr Ile Pro Lys Leu Asp 210 215 220

Thr Pro Asp Leu Leu Glu Val Gly Thr Gln Gln Lys Leu Phe Cys Ser 225 230 235 240

Leu Glu Ala Leu Phe Pro Ala Ser Glu Ala Arg Ile Tyr Leu Glu Leu 250 Gly Gln Met Pro Thr Gln Glu Ser Thr Asn Ser Ser Asp Ser Val 265 Ser Ala Thr Ala Leu Val Glu Val Thr Glu Glu Phe Asp Arg Thr Leu 275 Pro Leu Arg Cys Val Leu Glu Leu Ala Asp Gln Ile Leu Glu Thr Gln 295 Arg Thr Leu Thr Val Tyr Asn Phe Ser Ala Pro Val Leu Thr Leu Ser 310 315 Gln Leu Glu Val Ser Glu Gly Ser Gln Val Thr Val Lys Cys Glu Ala 330 His Ser Gly Ser Lys Val Val Leu Leu Ser Gly Val Glu Pro Arg Pro Pro Thr Pro Gln Val Gln Phe Thr Leu Asn Ala Ser Ser Glu Asp His 360 Lys Arg Ser Phe Phe Cys Ser Ala Ala Leu Glu Val Ala Gly Lys Phe 375 Leu Phe Lys Asn Gln Thr Leu Glu Leu His Val Leu Tyr Gly Pro Arg 390 395 Leu Asp Glu Thr Asp Cys Leu Gly Asn Trp Thr Trp Gln Glu Gly Ser Gln Gln Thr Leu Lys Cys Gln Ala Trp Gly Asn Pro Ser Pro Lys Met Thr Cys Arg Arg Lys Ala Asp Gly Ala Leu Leu Pro Ile Gly Val Val Lys Ser Val Lys Gln Glu Met Asn Gly Thr Tyr Val Cys His Ala Phe Ser Ser His Gly Asn Val Thr Arg Asn Val Tyr Leu Thr Val Leu Tyr 470 475 His Ser Gln Asn Asn Trp Thr Ile Ile Ile Leu Val Pro Val Leu Leu Val Ile Val Gly Leu Val Met Ala Ala Ser Tyr Val Tyr Asn Arg Gln Arg Lys Ile Arg Ile Tyr Lys Leu Gln Lys Ala Gln Glu Glu Ala Ile 515 Lys Leu Lys Gly Gln Ala Pro Pro

<210> 72

<211> 527

<212> PRT

<213> Cricetulus griseus

<400> 72

Met Ala Pro Thr Arg Ala Arg Pro Thr Pro Pro Leu Leu Leu Ala Leu 1 5 10 15

Val Ala Val Val Ile Pro Gly Pro Gly Ser Ala Gln Val Ser Ile His 20 25 30

Pro Lys Glu Ala Phe Leu Pro Arg Gly Ala Ser Met Gln Val Asn Cys 35 40 45

Ser Ser Ser Cys Ser Glu Asn Leu Ser Leu Gly Leu Glu Thr Gln Trp 50 55 60

Pro Lys Val Glu Leu Asp His Gly His Asn Trp Lys Leu Phe Glu Leu 65 70 75 80

Ser Asp Ile Gly Asp Asp Ser Lys Pro Leu Cys Phe Glu Asn Cys Gly 85 90 95

Pro Ile Gln Ser Ser Ala Ser Ala Thr Ile Val Leu Tyr Ser Phe Pro 100 105 110

Glu Arg Val Glu Leu Asp Arg Leu Pro Thr Trp Gln Pro Val Gly Lys 115 120 125

Asn Leu Thr Leu Arg Cys Leu Val Asp Gly Gly Thr Pro Arg Ser Gln 130 135 140

Leu Ser Val Lys Leu Leu Arg Gly Gly Glu Val Leu His Gln Glu Pro 145 150 155 160

Val Gly Val Asp Ser Arg Asn Pro Lys Glu Val Thr Val Thr Val Leu 165 170 175

Ala Ser Arg Asp Asp His Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu 180 185 190

Asp Leu Arg Pro Gln Gly Leu Ala Leu Phe Pro Asn Val Ser Val Ile 195 200 205

Arg Gln Leu Trp Thr Phe Asp Leu Pro Val Thr Glu Pro Lys Leu Asp 210 215 220

Thr Pro Asp Leu Leu Glu Val Gly Thr Val Gln Lys Val Met Cys Ser 225 230 235 240

Leu Gly Gly Leu Phe Pro Ala Ala Glu Ala Arg Ile Thr Leu Glu Leu 245 250 255

Gly Gly His Thr Leu Thr Ser Lys Ser Thr Asn His Arg Asp Leu Val 260 265 270 Ser Ala Thr Ala Leu Val Thr Ala Glu Met Glu Gly Thr Gln Gln Leu 275 280 285

Arg Cys Val Leu Glu Leu Ala Asp Gln Ile Leu Lys Ala Glu Arg Thr 290 295 300

Leu Ser Ile Tyr Asn Phe Ser Ala Pro Val Leu Thr Leu Ser Gln Gln 305 310 315 320

Glu Val Ser Glu Gly Ser Gln Val Thr Val Lys Cys Glu Ala Gln Gly 325 330 335

Gly Ala Gln Val Arg Leu Ser Gly Ala Pro Pro Gly Gln Val Gln Phe 340 345 350

Thr Leu Asn Ala Ser Ser Glu Asp His Glu Arg Ile Phe Thr Cys Ser 355 360 365

Ala Ala Leu Arg Val Ala Gly Gln Glu Leu Leu Lys Asn Gln Thr Leu 370 375 380

Lys Leu His Val Leu Tyr Gly Pro Arg Leu Asp Glu Asn Asp Cys Pro 385 390 395 400

Gly Asn Trp Thr Trp Pro Glu Gly Ser Gln Gln Asn Leu Ser Cys Gln 405 410 415

Ala Phe Gly Asn Pro Pro Pro Lys Leu Thr Cys Ser Arg Lys Thr Asp 420 425 430

Gly Ala Leu Leu Pro Ile Gly Glu Val Lys Thr Val Thr Trp Ala Met
435
440
445

Asn Gly Thr Tyr Val Cys His Ala Val Ser Ser His Gly Asn Ile Thr 450 455 460

Arg Glu Val Phe Leu Lys Val Leu Pro Lys Ser Pro Ile Trp Pro Ile 465 470 475 480

Ile Ile Ile Val Val Ile Leu Ala Thr Val Val Phe Val Gly Val Leu 485 490 495

Thr Ile Tyr Ile Tyr Asn Arg Gln Arg Lys Ile Arg Ile Tyr Lys Leu
500 505 510

Gln Arg Ala Gln Glu Glu Ala Met Lys Leu Lys Val Pro Pro His 515 520 525

<210> 73

<211> 544

<212> PRT

<213> Bos taurus

<400> 73

Met Ile Ala Ser Gly Pro Pro Pro Arg Val Tyr Trp Thr Ser Leu Ile 1 5 10 15 Phe Leu Leu Leu Ala Cys Cys Leu Leu Pro Thr Gly Ala Gln Gly Gln Thr Tyr Gln Val Arg Val Glu Pro Lys Asp Pro Val Val Pro Phe Gly Glu Pro Leu Val Val Asn Cys Thr Leu Asp Cys Pro Gly Pro Gly Leu Ile Ser Leu Glu Thr Ala Leu Ser Lys Glu Pro His Ser Arg Gly Leu Gly Trp Ala Ala Phe Arg Leu Thr Asn Val Thr Gly Asp Met Glu Ile Leu Cys Ser Gly Ile Cys Asn Lys Ser Gln Val Val Gly Phe Ser Asn Ile Thr Val Phe Gly Phe Pro Lys Arg Val Glu Leu Ala Pro Leu Pro 120 Leu Trp Gln Pro Val Gly Glu Glu Leu Asn Leu Ser Cys Leu Val Ser Gly Gly Ala Pro Arg Ala His Leu Ser Val Val Leu Leu Arg Gly Glu Glu Glu Leu Gly Arg Gln Pro Leu Gly Lys Glu Glu Pro Ala Lys Val Thr Phe Met Val Gln Pro Arg Arg Glu Asp His Gly Thr Asn Phe Ser 185 Cys Arg Ser Glu Leu Asp Leu Arg Ser Gln Gly Leu Glu Leu Phe Gln Asn Thr Ser Ala Pro Arg Lys Leu Gln Thr Tyr Ala Met Pro Lys Thr Ala Pro Arg Leu Val Phe Pro Arg Phe Trp Glu Met Glu Thr Ser Trp Pro Val Asn Cys Ser Leu Asn Gly Leu Phe Pro Ala Ser Glu Ala His 245 Ile Gln Leu Ala Leu Gly Asn Gln Met Leu Asn Ala Thr Val Val Ser 265 260 His Ala Asp Thr Leu Thr Ala Thr Ala Thr Ala Lys Thr Glu Glu 280 Gly Thr Gln Glu Ile Val Cys Asn Val Thr Leu Gly Val Glu Asn Arg 295 290 Glu Thr Arg Glu Ser Leu Val Ala Tyr Arg Phe Gln Gly Pro Asn Leu 315

Asn Leu Ser Glu Ser Asn Ala Thr Glu Gly Thr Pro Val Thr Val Thr Cys Ala Ala Gly Pro Gln Val Gln Val Met Leu Asp Gly Val Pro Ala 345 Ala Val Pro Gly Gln Pro Ala Gln Leu Gln Leu Lys Ala Thr Glu Met Asp Asp Arg Arg Thr Phe Phe Cys Asn Ala Thr Leu Lys Val His Gly 375 Val Thr Leu His Arg Asn Arg Ser Ile Gln Leu Arg Val Leu Tyr Gly Pro Thr Ile Asp Arg Ala Lys Cys Pro Gln Arg Leu Met Trp Lys Glu 410 Lys Thr Met His Ile Leu Gln Cys Gln Ala Arg Gly Asn Pro Asn Pro 420 Gln Leu Gln Cys Leu Arg Glu Gly Ser Lys Phe Lys Val Pro Val Gly Ile Pro Phe Leu Val Leu Leu Asn Tyr Ser Gly Thr Tyr Ser Cys Gln 455 Ala Ala Ser Ser Arg Gly Thr Asp Lys Met Leu Val Met Met Asp Val Gln Gly Arg Asn Pro Val Thr Ile Asn Ile Val Leu Gly Val Leu Ala Ile Leu Gly Leu Val Thr Leu Ala Ala Ala Ser Val Tyr Val Phe Trp

Ile Leu Gly Leu Val Thr Leu Ala Ala Ala Ser Val Tyr Val Phe Trp
500 505 510

Val Gln Arg Gln His Asp Ile Tyr His Leu Thr Pro Arg Ser Thr Arg 515 520 525

Trp Arg Leu Thr Ser Thr Gln Pro Val Thr Val Ala Glu Glu Leu Ser 530 535 540

<210> 74 <211> 537 <212> PRT

<213> Sus scrofa

<400> 74

Met Ala Pro Gly Ala Thr His Pro Gly Gln Leu Ala Leu Leu Ala Leu

1 5 10 15

Leu Leu Pro Leu Leu Gly Ala Leu Leu Pro Gly Leu Gly Gly Ala Glu 20 25 30

Ile Ser Met Trp Pro Leu Asn Thr Ile Ile Pro Lys Gly Gly Ser Met 35 40 45

Lys Val Asn Cys Ser Val Ala Cys Asp Gly Asn Ile Thr Ser Phe Gly Leu Glu Thr His Trp His Lys Thr Glu Val Asp His Arg Asp Lys Trp Lys Ile Phe Glu Leu Ser Asn Val Glu Asn Asp Gly Thr Leu Leu Cys His Ala Val Cys Gln Gly Asn Gln Thr Gln Val Gln Gly Asn Leu Thr Val Tyr Trp Phe Pro Glu Tyr Val Lys Leu Ala Asn Leu Ser Trp Gln 115 Arg Glu Gly Gln His Phe Asn Leu Ser Cys Gln Val Ser Gly Gly Ala 135 Pro Arg Thr Asn Leu Ser Ala Val Leu Phe Arg Gly Glu Glu Glu Leu 150 Phe Arg Gln Ser Val Gly Met Glu Glu Pro Ala Asn Val Thr Phe Arg 170 Met Leu Ala Ser Arq Lys Asp His Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu Asn Leu Gln Pro Gln Gly Leu Glu Leu Phe Trp Asn Ser Ser Ala Pro Leu Lys Leu Gln Thr Tyr Val Leu Pro Ala Thr His Pro His Leu Ala Thr Pro Glu Leu Val Glu Val Gly Thr Pro Val Ser Val Asn Cys Ser Leu Asp Gly Leu Phe Pro Ala Ser Glu Ala Thr Val His Leu 250 Ala Arg Gly Asp His Arg Pro Pro Leu Thr Ile Thr His Asn Gly Asp Ser Leu Leu Ala Lys Thr Trp Ile Asn Gly Thr Glu Lys Glu Gln Gly Thr Gln Tyr Leu Val Cys Glu Ile Met Leu Ala Asp Glu Lys Val Val 290 295 Thr Lys Lys Asn Val Thr Phe Tyr Ser Phe Pro Pro Pro Asn Leu Thr 310 315 Leu Ser Glu Pro Glu Val Ser Glu Gly Thr Thr Val Ser Ile Glu Cys 325 330 Gln Ala His Gly Glu Ala Val Val Thr Leu Asn Glu Val Pro Ala Glu

345

Pro Pro Ser Gln Arg Ala Gln Leu Lys Leu Asn Val Ser Ala Glu Asp 355 360 365

His Gly Arg Ser Phe Ser Cys Ser Ala Ala Leu Thr Val Ala Gly His 370 375 380

Val Leu Tyr Lys Asn Gln Thr Gln Val Leu Ser Val Leu Tyr Gly Pro 385 390 395 400

Arg Leu Asp Glu Arg Asp Cys Pro Gly Asn Trp Thr Trp Pro Glu Gly
405 410 415

Ser His Gln Thr Leu Thr Cys Gln Ala Arg Gly Asn Pro Thr Pro Lys 420 425 430

Leu Ile Cys Arg Arg Glu Gly Asp Gly Ala Leu Leu Pro Thr Gly Asp
435
440
445

Leu Gly Pro Val Lys Arg Glu Ile Thr Gly Thr Tyr Gln Cys Gln Ala 450 455 460

Thr Ser Ser Arg Gly Val Ala Thr Arg Val Val Val Val Asn Val Ile 465 470 475 480

His Asn Gln Asn Asn Met Val Ile Ile Ile Pro Val Ala Ala Val Ala 485 490 495

Ile Leu Gly Ser Val Gly Val Ala Ala Tyr Ile Tyr Asn Tyr Gln Arg
500 505 510

Lys Ile Gln Lys Tyr Glu Leu Gln Lys Ala Gln Glu Asn Ala Met 515 520 525

Lys Leu Ser Thr Pro Ala Ser Pro Pro 530 535

<210> 75

<211> 912

<212> PRT

<213> Oryctolagus cuniculus

<400> 75

Met Pro Gly Pro Ser Pro Gly Leu Arg Ala Leu Leu Gly Phe Trp Val

Ala Leu Gly Leu Gly Ile Leu Arg Leu Ser Ala Val Ala Gln Glu Pro 20 25 30

Phe Trp Ala Asp Leu Gln Pro Arg Val Ala Leu Val Glu Arg Gly Gly 35 40 45

Ser Leu Trp Leu Asn Cys Ser Thr Asn Cys Pro Arg Pro Glu Arg Gly
50 55 60

Gly Leu Glu Thr Ser Leu Arg Arg Asn Gly Pro Glu Gly Leu Arg Trp
65 70 75 80

Arg Ala Arg Gln Leu Val Asp Ile Arg Glu Pro Glu Thr Gln Pro Val Cys Phe Phe Arg Cys Ala Ala Thr Leu Gln Ala Arg Gly Leu Ile Arg Thr Phe Gln Arg Pro Asp Arg Val Glu Leu Val Pro Leu Pro Pro Trp Gln Pro Val Gly Glu Asn Phe Thr Leu Ser Cys Arg Val Pro Gly Ala 135 Gly Pro Arg Gly Ser Leu Thr Leu Thr Leu Leu Arg Gly Ala Gln Glu Leu Ile Arg Arg Ser Phe Ala Gly Glu Pro Ala Arg Ala Arg Gly Ala 165 170 Val Leu Thr Ala Thr Val Leu Ala Arg Arg Glu Asp His Gly Ala Asn Phe Ser Cys Arg Ala Glu Leu Asp Leu Arg Pro Gln Gly Leu Ala Leu Phe Glu Asn Ser Ser Ala Pro Arg Gln Leu Trp Thr Tyr Ala Leu Pro 215 Leu Asp Ser Pro Arg Leu Leu Ala Pro Arg Val Leu Glu Val Asp Ser Gln Ser Leu Val Ser Cys Thr Leu Asp Gly Leu Phe Pro Ala Ser Glu 245 250 Ala Gly Val His Leu Ala Leu Gly Asp Lys Arg Leu Asn Pro Glu Val Thr Leu Glu Gly Asp Ala Ile Val Ala Thr Ala Thr Ala Thr Ala Glu Glu Glu Gly Ile Lys Gln Leu Val Cys Ala Val Thr Leu Gly Gly Glu Arg Arg Glu Ser Arg Glu Asn Val Thr Val Tyr Ser Phe Pro Ala Pro 310 315 Leu Leu Thr Leu Ser Glu Pro Ser Ala Pro Glu Gly Lys Leu Val Thr 325 Val Thr Cys Thr Ala Gly Ala Arg Ala Leu Val Thr Leu Glu Gly Val 345 Pro Ala Ala Ala Pro Gly Gln Pro Ala Gln Leu Gln Phe Asn Ala Ser Glu Ser Asp Asp Gly Arg Ser Phe Phe Cys Asp Ala Thr Leu Glu Leu

Asp Gly Glu Thr Leu Ser Lys Asn Gly Ser Ala Glu Leu Arg Val Leu Tyr Ala Pro Arg Leu Asp Asp Ala Asp Cys Pro Arg Ser Trp Thr Trp 410 Pro Glu Gly Pro Glu Gln Thr Leu Arg Cys Glu Ala Arg Gly Asn Pro Thr Pro Ala Val His Cys Ala Arg Ser Asp Gly Gly Ala Val Leu Ala Leu Gly Leu Leu Gly Pro Val Thr Arg Ala Leu Ala Gly Thr Tyr Arg Cys Thr Ala Ala Asn Val Gln Gly Glu Ala Val Lys Asp Val Thr Leu Thr Val Glu Tyr Ala Pro Ala Leu Asp Ser Val Gly Cys Pro Glu Arg 485 Val Thr Trp Leu Glu Gly Thr Glu Ala Ser Leu Ser Cys Val Ala His Gly Val Pro Pro Pro Ser Val Ser Cys Val Arg Phe Arg Gln Ala Asp 520 Val Ile Glu Gly Leu Leu Leu Val Ala Arg Glu His Ala Gly Thr Tyr Arg Cys Glu Ala Ile Asn Ala Arg Ala Leu Ala Lys Asn Val Ala Val Thr Val Glu Tyr Gly Pro Ser Phe Glu Glu Arg Ser Cys Pro Ser Asn Trp Thr Trp Val Glu Gly Ser Glu Gln Leu Phe Ser Cys Glu Val Glu Gly Lys Pro Gln Pro Ser Val Gln Cys Val Gly Ser Glu Gly Ala Ser Glu Gly Leu Leu Pro Leu Ala Pro Leu Asn Pro Ser Pro Ser Asp Pro Ser Val Pro Arg Asp Leu Ala Pro Gly Ile Tyr Val Cys Asn Ala Thr Asn Pro Leu Gly Ser Ala Val Lys Thr Val Val Val Ser Ala Glu 650 645 Ser Pro Pro Gln Met Asp Asp Ser Thr Cys Pro Ser Asp Gln Thr Trp 660 Leu Glu Gly Ala Glu Ala Ala Gly Pro Ala Cys Ala Arg Gly Arg Pro 680 675

Ser Pro Arg Val Arg Cys Ser Arg Glu Gly Ala Pro Arg Pro Ala Arg 695 Pro Arg Val Ser Arg Glu Asp Ala Gly Thr Tyr Leu Cys Val Ala Thr 715 710 Asn Ala His Gly Ser Asp Ser Arg Thr Val Thr Val Gly Val Glu Tyr Arg Pro Val Val Ala Glu Leu Ala Ala Ser Pro Ser Gly Gly Val Arg 745 Pro Gly Gly Asn Phe Thr Leu Thr Cys Arg Ala Glu Ala Trp Pro Pro Ala Gln Ile Ser Trp Arg Ala Pro Pro Gly Ala Pro Asn Ile Gly Leu Ser Ser Asn Asn Ser Thr Leu Ser Val Pro Gly Ala Met Gly Ser His 790 Gly Gly Glu Tyr Glu Cys Glu Ala Thr Asn Ala His Gly His Ala Arg 810 Arg Ile Thr Val Arg Val Ala Gly Pro Trp Leu Trp Ile Ala Val Gly 825 Gly Ala Val Gly Gly Ala Val Leu Leu Ala Ala Gly Ala Gly Leu Ala Phe Tyr Val Gln Ser Thr Ala Cys Lys Lys Gly Glu Tyr Asn Val Gln

Glu Ala Glu Ser Ser Gly Glu Ala Val Cys Leu Asn Gly Ala Gly Gly 865 870 875 880

Gly Ala Gly Ser Gly Ala Glu Gly Gly Pro Glu Ala Glu Asp Ser Ala 885 890 895

Glu Ser Pro Ala Gly Gly Glu Val Phe Ala Ile Gln Leu Thr Ser Ala 900 905 910

<210> 76

<211> 917

<212> PRT

<213> Mus musculus

-400> 76

Met Pro Gly Pro Ser Pro Gly Leu Arg Arg Ala Leu Leu Gly Leu Trp

Ala Ala Leu Gly Leu Gly Ile Leu Gly Ile Ser Ala Val Ala Leu Glu 20 25 30

Pro Phe Trp Ala Asp Leu Gln Pro Arg Val Ala Leu Val Glu Pro Gly
35 40 45

Gly Ser Leu Trp Leu Asn Cys Ser Thr Asn Cys Pro Arg Pro Glu Arg Gly Gly Leu Glu Thr Ser Leu Arg Arg Asn Gly Thr Gln Arg Gly Leu Arg Trp Leu Ala Arg Gln Leu Val Asp Ile Arg Glu Pro Glu Thr Gln Pro Val Cys Phe Phe Arg Cys Ala Arg Arg Thr Leu Gln Ala Arg Gly 105 Leu Ile Arg Thr Phe Gln Arg Pro Asp Arg Val Glu Leu Val Pro Leu 115 Pro Ser Trp Gln Pro Val Gly Glu Asn Phe Thr Leu Ser Cys Arg Val 135 Pro Gly Ala Gly Pro Arg Ala Ser Leu Thr Leu Thr Leu Leu Arg Gly 150 Gly Gln Glu Leu Ile Arg Arg Ser Phe Val Gly Glu Pro Pro Arg Ala 170 Arg Gly Ala Met Leu Thr Ala Arg Val Leu Ala Arg Arg Glu Asp His Arg Val Asn Phe Ser Cys Leu Ala Glu Leu Asp Leu Arg Pro His Gly Leu Gly Leu Phe Ala Asn Ser Ser Ala Pro Arg Gln Leu Arg Thr Phe 215 Ala Met Pro Pro His Ser Pro Ser Leu Ile Ala Pro Arg Val Leu Glu Val Asp Ser Glu Arg Pro Val Thr Cys Thr Leu Asp Gly Leu Phe Pro Ala Pro Glu Ala Gly Val Tyr Leu Ser Leu Gly Asp Gln Arg Leu Asn Pro Asn Val Thr Leu Asp Gly Asp Ser Leu Val Ala Thr Ala Thr Ala Thr Ala Ser Ala Glu Gln Glu Gly Thr Lys Gln Leu Met Cys Val Val 295 290 Thr Leu Gly Gly Glu Thr Arg Glu Thr Gln Glu Asn Leu Thr Val Tyr 315 310 Ser Phe Pro Thr Pro Leu Leu Thr Leu Ser Glu Pro Glu Ala Pro Glu 325 Gly Lys Met Val Thr Ile Ser Cys Trp Ala Gly Ala Arg Ala Leu Val 345

Thr Leu Glu Gly Ile Pro Ala Ala Val Pro Gly Gln Pro Ala Glu Leu 360 Gln Leu Asn Val Thr Lys Asn Asp Asp Lys Arg Gly Phe Phe Cys Asp Ala Ala Leu Asp Val Asp Gly Glu Thr Leu Arg Lys Asn Gln Ser Ser Glu Leu Arg Val Leu Tyr Ala Pro Arg Leu Asp Asp Leu Asp Cys Pro 410 Arg Ser Trp Thr Trp Pro Glu Gly Pro Glu Gln Thr Leu His Cys Glu Ala Arq Gly Asn Pro Glu Pro Ser Val His Cys Ala Arg Pro Glu Gly Gly Ala Val Leu Ala Leu Gly Leu Leu Gly Pro Val Thr Arg Ala Leu Ala Gly Thr Tyr Arg Cys Thr Ala Val Asn Gly Gln Gly Gln Ala Val Lys Asp Val Thr Leu Thr Val Glu Tyr Ala Pro Ala Leu Asp Ser Val 485 Gly Cys Pro Glu His Ile Thr Trp Leu Glu Gly Thr Glu Ala Ser Leu 505 Ser Cys Val Ala Pro Gly Val Pro Pro Pro Ser Val Ser Cys Val Arg Ser Gly Lys Glu Glu Val Met Glu Gly Pro Leu Arg Val Ala Arg Glu His Ala Gly Thr Tyr Arg Cys Glu Ala Ile Asn Ala Arg Gly Ser Ala Ala Lys Asn Val Ala Val Thr Val Glu Tyr Gly Pro Ser Phe Glu Glu Leu Gly Cys Pro Ser Asn Trp Thr Trp Val Glu Gly Ser Gly Lys Leu 585 Phe Ser Cys Glu Val Asp Gly Lys Pro Glu Pro Arg Val Glu Cys Val Gly Ser Glu Gly Ala Ser Glu Gly Ile Val Leu Pro Leu Val Ser Ser Asn Ser Gly Pro Arg Asn Ser Met Thr Pro Gly Asn Leu Ser Pro Gly 630 625 Ile Tyr Leu Cys Asn Ala Thr Asn Arg His Gly Ser Thr Val Lys Thr

650

Val Val Ser Ala Glu Ser Pro Pro Gln Met Asp Glu Ser Ser Cys 660 665 670

Pro Ser His Gln Thr Trp Leu Glu Gly Ala Glu Ala Thr Ala Leu Ala 675 680 685

Cys Ser Ala Arg Gly Arg Pro Ser Pro Arg Val His Cys Ser Arg Glu 690 695 700

Gly Ala Ala Arg Leu Glu Arg Leu Gln Val Ser Arg Glu Asp Ala Gly 705 710 715 720

Thr Tyr Arg Cys Val Ala Thr Asn Ala His Gly Thr Asp Ser Arg Thr
725 730 735

Val Thr Val Gly Val Glu Tyr Arg Pro Val Val Ala Glu Leu Ala Ala
740 745 750

Ser Pro Pro Ser Val Arg Pro Gly Gly Asn Phe Thr Leu Thr Cys Arg
755 760 765

Ala Glu Ala Trp Pro Pro Ala Gln Ile Ser Trp Arg Ala Pro Pro Gly 770 780

Ala Leu Asn Leu Gly Leu Ser Ser Asn Asn Ser Thr Leu Ser Val Ala 785 790 795 800

Gly Ala Met Gly Ser His Gly Gly Glu Tyr Glu Cys Ala Ala Thr Asn  $805 \\ 810 \\ 815$ 

Ala His Gly Arg His Ala Arg Arg Ile Thr Val Arg Val Ala Gly Pro 820 825 830

Trp Leu Trp Val Ala Val Gly Gly Ala Ala Gly Gly Ala Ala Leu Leu 835 840 845

Ala Ala Gly Ala Gly Leu Ala Phe Tyr Val Gln Ser Thr Ala Cys Lys 850 860

Lys Gly Glu Tyr Asn Val Gln Glu Ala Glu Ser Ser Gly Glu Ala Val 865 870 875 880

Cys Leu Asn Gly Ala Gly Gly Thr Pro Gly Ala Glu Gly Gly Ala Glu 885 890 895

Thr Pro Gly Thr Ala Glu Ser Pro Ala Asp Gly Glu Val Phe Ala Ile 900 905 910

Gln Leu Thr Ser Ser 915

<210> 77

<211> 548

<212> PRT

<213> Mus musculus

<400> 77

Met Lys Met Leu Leu Leu Gly Val Trp Thr Leu Leu Ala Leu Ile Pro 1 5 10 15

Cys Pro Gly Ala Ala Glu Glu Leu Phe Gln Val Ser Val His Pro Asn 20 25 30

Glu Ala Leu Val Glu Phe Gly His Ser Leu Thr Val Asn Cys Ser Thr 35 40 45

Thr Cys Pro Asp Pro Gly Pro Ser Gly Ile Glu Thr Phe Leu Lys Lys 50 55 60

Thr Gln Leu Ser Lys Gly Ser Gln Trp Lys Glu Phe Leu Leu Glu Asp
65 70 75 80

Ile Thr Glu Asp Leu Val Leu Gln Cys Phe Phe Ser Cys Ala Gly Glu
85 90 95

Gln Lys Asp Thr Val Leu Ala Ile Thr Met Tyr Gln Pro Pro Glu Gln
100 105 110

Val Ile Leu Asp Leu Gln Pro Glu Trp Val Ala Val Asp Glu Ala Phe 115 120 125

Thr Val Thr Cys His Val Pro Ser Val Ala Pro Leu Gln Ser Leu Thr 130 135 140

Leu Thr Leu Leu Gln Gly Asp Gln Glu Leu His Arg Lys Asp Phe Leu 145 150 155 160

Ser Leu Ser Leu Val Ser Gln Arg Ala Glu Val Thr Ala Thr Val Arg 165 170 175

Ala His Arg Asp Asn Asp Arg Arg Asn Phe Ser Cys Arg Ala Glu Leu 180 185 190

Asp Leu Ser Pro His Gly Gly Gly Leu Phe His Gly Ser Ser Ala Thr 195 200 205

Lys Gln Leu Arg Ile Phe Glu Phe Ser Gln Asn Pro Gln Ile Trp Val 210 215 220

Pro Ser Leu Leu Glu Val Gly Lys Ala Glu Ile Val Ser Cys Glu Val 225 230 235 240

Thr Arg Val Phe Pro Ala Gln Glu Ala Val Phe Arg Met Phe Leu Glu 245 250 255

Asp Gln Glu Leu Ser Pro Phe Ser Ser Trp Arg Glu Asp Ala Ala Trp 260 265 270

Ala Ser Ala Thr Ile Gln Ala Met Glu Thr Gly Asp Gln Glu Leu Thr 275 280 285

Cys Leu Val Ser Leu Gly Pro Val Glu Gln Lys Thr Arg Lys Pro Val 290 295 300 Tyr Val Tyr Ser Phe Pro Pro Pro Ile Leu Glu Ile Glu Asp Ala Tyr 315 310 Pro Leu Ala Gly Thr Asp Val Asn Val Thr Cys Ser Gly His Val Leu 330 325 Thr Ser Pro Ser Pro Thr Leu Arg Leu Gln Gly Ser Leu Asn His Ser Ala Pro Gly Lys Pro Ala Trp Leu Leu Phe Thr Ala Arg Glu Glu Asp 360 Asp Gly Arg Thr Leu Ser Cys Glu Ala Ser Leu Glu Val Gln Gly Gln Arg Leu Val Arg Thr Thr Glu Ser Gln Leu His Val Leu Tyr Lys Pro Arg Phe Gln Glu Ser Arg Cys Pro Gly Asn Gln Ile Trp Val Glu Gly 405 Met His Gln Met Leu Ala Cys Ile Pro Glu Gly Asn Pro Thr Pro Val Leu Val Cys Val Trp Asn Gly Met Ile Phe Asp Leu Asp Val Pro Gln 440 Lys Ala Thr Gln Asn His Thr Gly Thr Tyr Cys Cys Thr Ala Thr Asn Pro Leu Gly Ser Val Ser Lys Asp Ile Thr Ile Ile Val Gln Gly Leu 470 Pro Glu Gly Ile Ser Ser Ser Thr Ile Phe Ile Ile Ile Phe Thr Leu Gly Met Ala Val Ile Thr Val Ala Leu Tyr Leu Asn Tyr Gln Pro 505 Cys Lys Gly Asn Ser Arg Lys Arg Met His Arg Pro Arg Glu Gln Ser Lys Gly Glu Glu Ser Gln Phe Ser Asp Ile Arg Ala Glu Glu Cys His Ala His Leu Cys 545

<210> 78

<211> 548

<212> PRT

<213> Rattus norvegicus

<400> 78

Met Lys Met Leu Leu Gly Ile Trp Thr Leu Leu Ala Leu Ile Pro 1 5 10 15 Cys Pro Gly Thr Thr Glu Val Leu Phe Gln Val Ser Val His Pro Asn Gln Ala Leu Val Glu Phe Gly His Ser Leu Thr Ile Asn Cys Ser Thr Thr Cys Pro Asp Pro Gly Pro Ser Gly Ile Glu Thr Phe Leu Lys Lys Thr Gln Leu Ser Lys Gly Ser Gln Trp Lys Glu Phe Leu Leu Glu Gly Ile Thr Glu Asn Ser Val Leu Gln Cys Phe Phe Ser Cys Ala Gly Val Gln Lys Asp Thr Ala Leu Asp Ile Thr Met Tyr Gln Pro Pro Glu Gln Val Ile Leu Asp Leu Gln Pro Glu Trp Val Ala Ile Asp Glu Ala Phe 115 Thr Val Lys Cys His Val Pro Ser Val Ala Pro Leu Gln Ser Leu Thr 135 Leu Thr Leu Leu Gln Gly Asp Gln Glu Leu His Arg Lys Asp Phe Leu 150 Ser Leu Ser Leu Val Ser Gln Arg Ala Glu Val Thr Val Asn Val Arg 170 Ala Gln Arg Glu Asn Asp Arg His Asn Phe Ser Cys Arg Ala Glu Leu Asp Leu Ser Pro His Gly Gly Gly Leu Phe His Gly Ser Ser Ala Thr Lys Gln Leu Arg Ile Phe Glu Phe Ser Gln Asn Pro Gln Ile Leu Val 215 Pro Ser Leu Leu Glu Val Gly Met Ala Glu Thr Met Ser Cys Glu Val Val Arg Val Phe Pro Ala Gln Glu Ala Val Phe Arg Met Phe Leu Glu 250 Asp Gln Glu Leu Ser Pro Phe Ser Ser Trp Lys Gly Asp Ala Ala Trp Ala Ser Ala Thr Ile Gln Ala Met Glu Thr Gly Asp Gln Glu Leu Thr

Cys Leu Val Ser Val Gly Pro Val Glu Gln Lys Ala Arg Lys Pro Val

His Val Tyr Ser Phe Pro Pro Pro Val Leu Glu Ile Glu Asp Ala Tyr

315

295

310

Pro Gln Ala Gly Thr Asp Val Asn Val Thr Cys Ser Gly His Val Leu 325 330 335

Thr Ser Pro Ser Pro Thr Leu Arg Leu Gln Gly Ser Leu Asn Leu Ser 340 345 350

Ala Pro Gly Glu Pro Ala Trp Leu Arg Phe Thr Ala Arg Glu Glu Asp 355 360 365

Asp Gly Arg Thr Leu Ser Cys Glu Ala Ser Leu Val Val Gln Gly Gln 370 375 380

Arg Leu Val Lys Thr Thr Lys Ile Gln Leu His Val Leu Tyr Lys Pro 385 390 395 400

Arg Phe Gln Glu Ser Asp Cys Pro Gly Asn Gln Ile Trp Val Glu Gly 405 410 415

Met Asp Gln Met Leu Ala Cys Ile Pro Glu Gly Asn Pro Ile Pro Ala 420 425 430

Leu Val Cys Ile Trp Asn Gly Met Thr Phe Asp Leu Glu Val Pro Gln 435 440 445

Lys Ala Thr Gln Asn His Thr Gly Thr Tyr Ser Cys Thr Ala Thr Asn 450 455 460

Ser Leu Gly Ser Val Ser Lys Asp Ile Ala Val Leu Val Gln Gly Leu 465 470 475 480

His Glu Gly Ile Ser Ser Ser Thr Ile Phe Ile Ile Ile Ile Phe Thr 485 490 495

Leu Gly Met Ala Val Ile Thr Ile Ala Leu Tyr Leu Asn Tyr Gln Pro 500 505 510

Cys Lys Arg Asn Gly Arg Lys Arg Thr His Arg Gln Lys Glu Gln Asn 515 520 525

Lys Gly Glu Glu Arg Gln Phe Ser Asp Ile Gln Ala Glu Glu Cys His 530 535 540

Ala His Leu Cys 545

<210> 79

<211> 396

<212> PRT

<213> Mus musculus

<400> 79

Met Gly Ala Pro Ser Ala Leu Pro Leu Leu Leu Leu Leu Ala Cys Ser

Trp Ala Pro Gly Gly Ala Asn Leu Ser Gln Asp Asp Ser Gln Pro Trp 20 25 30

Thr Ser Asp Glu Thr Val Val Ala Gly Gly Thr Val Val Leu Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu Gln Trp Ser Asn Pro Ala Gln Gln Thr Leu Tyr Phe Gly Glu Lys Arg Ala Leu Arg Asp Asn Arg Ile Gln Leu Val Ser Ser Thr Pro His Glu Leu Ser Ile Ser Ile Ser Asn Val Ala Leu Ala Asp Glu Gly Glu Tyr Thr Cys Ser Ile Phe Thr Met Pro Val Arg Thr Ala Lys Ser Leu Val Thr Val Leu Gly Ile Pro Gln Lys Pro Ile Ile Thr Gly Tyr Lys Ser Ser Leu Arg Glu Lys Glu 135 Thr Ala Thr Leu Asn Cys Gln Ser Ser Gly Ser Lys Pro Ala Ala Gln 150 Leu Thr Trp Arg Lys Gly Asp Gln Glu Leu His Gly Asp Gln Thr Arg Ile Gln Glu Asp Pro Asn Gly Lys Thr Phe Thr Val Ser Ser Ser Val Ser Phe Gln Val Thr Arg Glu Asp Asp Gly Ala Asn Ile Val Cys Ser Val Asn His Glu Ser Leu Lys Gly Ala Asp Arg Ser Thr Ser Gln Arg Ile Glu Val Leu Tyr Thr Pro Thr Ala Met Ile Arg Pro Glu Pro Ala 230 His Pro Arg Glu Gly Gln Lys Leu Leu His Cys Glu Gly Arg Gly Asn Pro Val Pro Gln Gln Tyr Val Trp Val Lys Glu Gly Ser Glu Pro Pro Leu Lys Met Thr Gln Glu Ser Ala Leu Ile Phe Pro Phe Leu Asn 275 Lys Ser Asp Ser Gly Thr Tyr Gly Cys Thr Ala Thr Ser Asn Met Gly 295 Ser Tyr Thr Ala Tyr Phe Thr Leu Asn Val Asn Asp Pro Ser Pro Val 305 310 Pro Ser Ser Ser Thr Tyr His Ala Ile Ile Gly Gly Ile Val Ala

330

Phe Ile Val Phe Leu Leu Leu Ile Leu Leu Ile Phe Leu Gly His Tyr 340 345 350

Leu Ile Arg His Lys Gly Thr Tyr Leu Thr His Glu Ala Lys Gly Ser 355 360 365

Asp Asp Ala Pro Asp Ala Asp Thr Ala Ile Ile Asn Ala Glu Gly Gly 370 375 380

Gln Ser Gly Gly Asp Asp Lys Lys Glu Tyr Phe Ile 385 390 395

<210> 80

<211> 662

<212> PRT

<213> Homo sapiens

<400> 80

Met Glu Ser Lys Thr Glu Lys Trp Met Glu Arg Ile His Leu Asn Val 1 5 10 15

Ser Glu Arg Pro Phe Pro Pro His Ile Gln Leu Pro Pro Glu Ile Gln 20 25 30

Glu Ser Gln Glu Val Thr Leu Thr Cys Leu Leu Asn Phe Ser Cys Tyr 40 45

Gly Tyr Pro Ile Gln Leu Gln Trp Leu Leu Glu Gly Val Pro Met Arg 50 55 60

Gln Ala Ala Val Thr Ser Thr Ser Leu Thr Ile Lys Ser Val Phe Thr 65 70 75 80

Arg Ser Glu Leu Lys Phe Ser Pro Gln Trp Ser His His Gly Lys Ile 85 90 95

Val Thr Cys Gln Leu Gln Asp Ala Asp Gly Lys Phe Leu Ser Asn Asp 100 105 110

Thr Val Gln Leu Asn Val Lys His Thr Pro Lys Leu Glu Ile Lys Val 115 120 125

Thr Pro Ser Asp Ala Ile Val Arg Glu Gly Asp Ser Val Thr Met Thr 130 135 140

Cys Glu Val Ser Ser Thr Asn Pro Glu Tyr Thr Thr Val Ser Trp Leu 145 150 155 160

Lys Asp Gly Thr Ser Leu Lys Lys Gln Asn Thr Phe Thr Leu Asn Leu 165 170 175

Arg Glu Val Thr Lys Asp Gln Ser Gly Lys Tyr Cys Cys Gln Val Ser 180 185 190

Asn Asp Val Gly Pro Gly Arg Ser Glu Glu Val Phe Leu Gln Val Gln
195 200 205

Tyr Ala Pro Glu Pro Ser Thr Val Gln Ile Leu His Ser Pro Ala Val 215 Glu Gly Ser Gln Val Glu Phe Leu Cys Met Ser Leu Ala Asn Pro Leu 235 Pro Thr Asn Tyr Thr Trp Tyr His Asn Gly Lys Glu Met Gln Gly Arg Thr Glu Glu Lys Val His Ile Pro Lys Ile Leu Pro Trp His Ala Gly 265 Thr Tyr Ser Cys Val Ala Glu Asn Ile Leu Gly Thr Gly Gln Arg Gly Pro Gly Ala Glu Leu Asp Val Gln Tyr Pro Pro Lys Lys Val Thr Thr 295 Val Ile Gln Asn Pro Met Pro Ile Arg Glu Gly Asp Thr Val Thr Leu Ser Cys Asn Tyr Asn Ser Ser Asn Pro Ser Val Thr Arg Tyr Glu Trp 330 Lys Pro His Gly Ala Trp Glu Glu Pro Ser Leu Gly Val Leu Lys Ile Gln Lys Val Gly Trp Asp Asn Thr Thr Ile Ala Cys Ala Arg Cys Asn Ser Trp Cys Ser Trp Ala Ser Pro Val Ala Leu Asn Val Gln Tyr Ala 375 Pro Arg Asp Val Arg Val Arg Lys Ile Lys Pro Leu Ser Glu Ile His Ser Gly Asn Ser Val Ser Leu Gln Cys Asp Phe Ser Ser His Pro Lys Glu Val Gln Phe Phe Trp Glu Lys Asn Gly Arg Leu Leu Gly Lys Glu Ser Gln Leu Asn Phe Asp Ser Ile Ser Pro Glu Asp Ala Gly Ser Tyr Ser Cys Trp Val Asn Asn Ser Ile Gly Gln Thr Ala Ser Lys Ala Trp Thr Leu Glu Val Leu Tyr Ala Pro Arg Arg Leu Arg Val Ser Met 475 Ser Pro Gly Asp Gln Val Met Glu Gly Lys Ser Ala Thr Leu Arg Cys 485 Glu Ser Asp Ala Asn Pro Pro Val Ser His Tyr Thr Trp Phe Asp Trp 505

Asn Asn Gln Ser Leu Pro Tyr His Ser Gln Lys Leu Arg Leu Glu Pro 515 520 525

Val Lys Val Gln His Ser Gly Ala Tyr Trp Cys Gln Gly Thr Asn Ser 530 535 540

Val Gly Lys Gly Arg Ser Pro Leu Ser Thr Leu Thr Val Tyr Tyr Ser 545 550 555 560

Pro Glu Thr Ile Gly Arg Arg Val Ala Val Gly Leu Gly Ser Cys Leu
565 570 575

Ala Ile Leu Ile Leu Ala Ile Cys Gly Leu Lys Leu Gln Arg Arg Asp 580 585 590

Ala Glu Ser Ser Glu Met Gln Arg Pro Pro Arg Thr Cys Asp Asp Thr
595 600 605

Val Thr Tyr Ser Ala Leu His Lys Arg Gln Val Gly Asp Tyr Glu Asn 610 620

Val Ile Pro Asp Phe Pro Glu Asp Glu Gly Ile His Tyr Ser Glu Leu 625 630 635 640

Ile Gln Phe Gly Val Gly Glu Arg Pro Gln Ala Gln Glu Asn Val Asp 645 650 655

Tyr Val Ile Leu Lys His 660

<210> 81

<211> 505

<212> PRT

<213> Pan troglodytes

<400> 81

Gln Thr Ser Val Ser Pro Pro Lys Val Ile Leu Pro Arg Gly Gly Ser

1 5 10 15

Val Gln Val Thr Cys Ser Thr Ser Cys Asp Gln Pro Asp Leu Leu Gly
20 25 30

Ile Glu Thr Pro Leu Pro Lys Lys Glu Leu Leu Gly Gly Asn Asn 35 40 45

Trp Lys Val Tyr Glu Leu Ser Asn Val Gln Glu Asp Ser Gln Pro Met 50 55 60

Cys Tyr Ser Asn Cys Pro Asp Gly Gln Ser Thr Ala Lys Thr Phe Leu 65 70 75 80

Thr Val Tyr Trp Thr Pro Glu Arg Val Glu Leu Ala Pro Leu Pro Ser 85 90 95

Trp Gln Pro Val Gly Lys Asp Leu Thr Leu Arg Cys Gln Val Glu Gly
100 105 110

Gly Ala Pro Arg Ala Asn Leu Thr Val Val Leu Leu Arg Gly Glu Lys 120 Glu Leu Lys Arg Glu Pro Ala Val Gly Glu Pro Ala Glu Val Thr Thr 140 Thr Val Leu Val Glu Arg Asp His His Gly Ala Asn Phe Ser Cys Arg Thr Glu Leu Asp Leu Arg Pro Gln Gly Leu Gln Leu Phe Glu Asn Thr 170 Ser Ala Pro His Gln Leu Gln Thr Phe Val Leu Pro Ala Thr Pro Pro 180 Gln Leu Val Ser Pro Arg Val Leu Glu Val Asp Thr Gln Gly Thr Val 200 Val Cys Ser Leu Asp Gly Leu Phe Pro Val Leu Glu Ala Gln Val His Leu Ala Leu Gly Asp Gln Arg Leu Asn Pro Thr Val Thr Tyr Gly Asn Asp Ser Phe Ser Ala Lys Ala Ser Val Ser Val Thr Ala Glu Asp Glu 245 Gly Thr Gln Arg Leu Thr Cys Ala Val Ile Leu Gly Asn Gln Ser Arg Glu Thr Leu Gln Thr Val Thr Ile Tyr Ser Phe Pro Ala Pro Asn Val 280 Ile Leu Thr Lys Pro Glu Val Ser Glu Gly Thr Glu Val Thr Val Lys Cys Glu Ala His Pro Arg Ala Lys Val Thr Leu Asn Gly Val Pro Ala Gln Pro Val Gly Pro Arg Val Gln Leu Leu Leu Lys Ala Thr Pro Glu Asp Asn Gly Arg Ser Phe Ser Cys Ser Ala Thr Leu Glu Val Ala Gly 345 Gln Leu Ile His Lys Asn Gln Thr Arg Glu Leu Arg Val Leu Tyr Gly Pro Arg Leu Asp Glu Arg Asp Cys Pro Gly Asn Trp Thr Trp Pro Glu 375 Asn Ser Gln Gln Thr Pro Met Cys Gln Ala Ser Gly Asn Pro Leu Pro Glu Leu Lys Cys Leu Lys Asp Gly Thr Phe Pro Leu Pro Val Gly Glu Ser Val Thr Val Thr Arg Asp Leu Glu Gly Thr Tyr Leu Cys Arg Ala 420 425 430

Arg Ser Thr Gln Gly Glu Val Thr Arg Lys Val Thr Val Asn Val Leu 435 440 445

Ser Pro Arg Tyr Glu Ile Val Ile Ile Thr Val Val Ala Ala Ala Val 450 455 460

Ile Met Gly Thr Ala Gly Leu Ser Thr Tyr Leu Tyr Asn Arg Gln Arg 465 470 475 480

Lys Ile Arg Lys Tyr Arg Leu Gln Gln Ala Gln Lys Gly Thr Pro Met 485 490 495

Lys Pro Asn Thr Gln Ala Thr Pro Pro 500 505

<210> 82

<211> 447

<212> PRT

<213> Mus musculus

<220>

<221> MOD RES

<222> (12)

<223> Any amino acid

<220>

<221> MOD RES

<222> (77)..(80)

<223> Any amino acid

<220>

<221> MOD\_RES

 $<222> (14\overline{5})..(147)$ 

<223> Any amino acid

<220>

<221> MOD\_RES

<222> (155)..(163)

<223> Any amino acid

<220>

<221> MOD RES

<222> (268)..(269)

<223> Any amino acid

<220>

<221> MOD\_RES

<222> (279)

<223> Any amino acid

<220>

<221> MOD RES

<222> (282)

<223> Any amino acid

<400> 82

Glu Asp Ser Gln Pro Met Cys Tyr Ser Asn Cys Xaa Asp Gly Gln Ser

Thr Ala Lys Thr Phe Leu Thr Val Tyr Trp Thr Pro Glu Arg Val Glu
20 25 30

Leu Ala Pro Leu Pro Ser Trp Gln Pro Val Gly Lys Asn Leu Thr Leu 35 40 45

Arg Cys Gln Val Glu Gly Gly Ala Pro Arg Ala Asn Leu Thr Val Val 50 55 60

Leu Leu Arg Gly Glu Lys Glu Leu Lys Arg Glu Pro Xaa Xaa Xaa 65 70 75 80

Pro Ala Glu Val Thr Thr Thr Val Leu Val Arg Arg Asp His His Gly 85 90 95

Ala Asn Phe Ser Cys Arg Thr Glu Leu Asp Leu Arg Pro Gln Gly Leu 100 105 110

Glu Leu Phe Glu Asn Thr Ser Ala Pro Tyr Gln Leu Gln Thr Phe Val 115 120 125

Leu Pro Ala Thr Pro Pro Gln Leu Val Ser Pro Arg Val Leu Glu Val 130 135 140

Xaa Xaa Xaa Gly Thr Val Val Cys Ser Leu Xaa Xaa Xaa Xaa Xaa 145 150 160

Xaa Xaa Xaa Gln Val His Leu Ala Leu Gly Asp Gln Arg Leu Asn Pro 165 170 175

Thr Val Thr Tyr Gly Asn Asp Ser Phe Ser Ala Lys Ala Ser Val Ser 180 185 190

Val Thr Ala Glu Asp Glu Gly Thr Gln Arg Leu Thr Cys Ala Val Ile 195 200 205

Leu Gly Asn Gln Ser Gln Glu Thr Leu Gln Thr Val Thr Ile Tyr Ser 210 215 220

Phe Pro Ala Pro Asn Val Ile Leu Thr Lys Pro Glu Val Ser Glu Gly 225 230 235 240

Thr Glu Val Thr Val Lys Cys Glu Ala His Pro Arg Ala Lys Val Thr
245 250 255

Leu Asn Gly Val Pro Ala Gln Pro Leu Gly Pro Xaa Xaa Gln Leu Leu 260 265 270

Leu Lys Ala Thr Pro Glu Xaa Asn Gly Xaa Ser Phe Ser Cys Ser Ala 275 280 285

Thr Leu Glu Val Ala Gly Gln Leu Ile His Lys Asn Gln Thr Arg Glu 290 295 300 Leu Arg Val Leu Tyr Gly Pro Arg Leu Asp Glu Arg Asp Cys Pro Gly 305 310 315 320

Asn Trp Thr Trp Pro Glu Asn Ser Gln Gln Thr Pro Met Cys Gln Ala 325 330 335

Trp Gly Asn Pro Leu Pro Glu Leu Lys Cys Leu Lys Asp Gly Thr Phe 340 345 350

Pro Leu Pro Ile Gly Glu Ser Val Thr Val Thr Arg Asp Leu Glu Gly 355 360 365

Thr Tyr Leu Cys Arg Ala Arg Ser Thr Gln Gly Glu Val Thr Arg Glu 370 380

Val Thr Val Asn Val Leu Ser Pro Arg Tyr Glu Ile Val Ile Ile Thr 385 390 395 400

Val Val Ala Ala Ala Val Ile Met Gly Thr Ala Gly Leu Ser Thr Tyr 405 410 415

Leu Tyr Asn Arg Gln Arg Lys Ile Lys Lys Tyr Arg Leu Gln Gln Ala 420 425 430

Gln Lys Gly Thr Pro Met Lys Pro Asn Thr Gln Ala Thr Pro Pro 435 440 445

<210> 83

<211> 528

<212> PRT

<213> Canis familiaris

<400> 83

Ala Pro Ala Leu Pro Arg Leu Pro Ala Leu Leu Ala Leu Leu Gly Ala 1 5 10 15

Leu Leu Pro Gly Leu Gly Gly Ala Gln Thr Ser Val Asp Pro Ala Glu 20 25 30

Ala Ile Ile Leu Arg Gly Gly Ser Val Gln Val Asn Cys Ser Thr Ser 35 40 45

Cys Asn Gln Thr Ser Ile Phe Gly Leu Glu Thr Leu Leu Thr Lys Thr 50 55 60

Glu Val Thr Ser Gly Asp Asn Trp Val Leu Phe Glu Leu Thr Asp Val
65 70 75 80

Gln Glu Asp Ser Lys Leu Ile Cys Phe Ser Asn Cys His Asp Glu Thr 85 90 95

Met Ala Pro Ile Asp Leu Thr Val Tyr Trp Phe Pro Glu Arg Val Glu 100 105 110

Leu Ala Pro Leu Pro Arg Trp Gln Pro Val Gly Glu Asn Leu Thr Met 115 120 125

Thr Cys Gln Val Ala Gly Gly Ala Pro Arg Thr Asn Leu Thr Val Val Leu Leu Arg Gly Glu Glu Leu Ser Arg Gln Pro Ala Val Gly Glu Pro Ala Glu Val Thr Phe Thr Val Ala Val Gly Arg Glu Asp His Leu 170 Ala Asn Phe Ser Cys Arg Thr Asp Leu Asp Leu Arg His Arg Gly Leu Gly Leu Phe Gln Asn Ser Ser Ala Pro Arg Gln Leu Gln Thr Phe Val Leu Pro Glu Thr Pro Pro Arg Leu Ala Thr Pro Pro Ile Val Glu Val 215 Gly Thr Gln Trp Ser Val Asp Cys Thr Met Asp Gly Val Phe Pro Ala Ser Glu Ala Gln Val His Leu Ala Leu Ala Glu Glu Arg Leu His Ser 245 Thr Val Leu Tyr Lys Lys Asp Ser Leu Leu Ala Thr Ala Asn Val Lys Ala Asn Pro Glu Asp Glu Gly Thr Gln Gln Leu Trp Cys Glu Val Thr Leu Gly Asp Glu Asn Arg Arg Trp Gln Glu Asn Val Thr Phe Tyr Ser Phe Pro Ala Pro Asn Leu Thr Leu Ser Glu Pro Glu Val Ser Glu Trp 310 Thr Thr Val Thr Val Glu Cys Glu Ala Pro Ala Gly Val Val Ser Leu Ser Gly Leu Pro Ser Gly Leu Ala Val Pro Arg Ala Gln Phe Gln Leu Asn Ala Ser Ala Ala Asp Asn Arg Arg Ser Phe Ser Cys Ser Ala Ala Leu Glu Val Ala Gly His Met Leu Gln Lys Asn Gln Thr Arg Glu 375 Leu His Val Leu Tyr Gly Pro Arg Leu Asp Gln Arg Asp Cys Pro Gly Asn Trp Thr Trp Glu Glu Gly Phe His Gln Thr Leu Lys Cys Gln Ala 410 405 Trp Gly Asn Pro Val Pro Glu Leu Lys Cys His Arg Lys Gly Asp Asp 425 420

Ala Leu Leu Pro Ile Gly Asp Leu Arg Pro Val Lys Arg Glu Val Ala 435 440 445

Gly Thr Tyr Leu Cys Gln Ala Arg Ser Pro Arg Gly Glu Ile Thr Arg 450 450 460

Glu Val Val Ile Asn Val Ile Tyr His Gln Asn Asn Ile Leu Ile Ile 465 470 475 480

Ile Leu Val Thr Thr Ile Val Ile Leu Gly Thr Val Ser Val Ala Ala 485 490 495

Tyr Leu Tyr Asn Arg Gln Arg Lys Ile Gln Lys Tyr Lys Leu Gln Lys
500 505 510

Ala Gln Glu Ala Ala Ala Met Lys Leu Asn Thr Pro Ala Thr Pro Pro 515 520 525

<210> 84

<211> 535

<212> PRT

<213> Bos taurus

<400> 84

Met Ala Leu Gly Ala Ala Pro Ala Ala Gln Leu Ala Leu Leu Ala Leu

1 10 15

Leu Gly Thr Leu Leu Pro Gly Pro Gly Gly Ala Gly Ile Ser Ile His
20 25 30

Pro Ser Lys Ala Ile Ile Pro Arg Gly Asp Ser Leu Thr Val Asn Cys
35 40 45

Ser Asn Ser Cys Asp Gln Lys Ser Thr Phe Gly Leu Glu Thr Val Leu 50 55 60

Ile Lys Glu Glu Val Gly Arg Gly Asp Asn Trp Lys Val Phe Gln Leu 65 70 75 80

Arg Asp Val Glu Glu Asp Ile Glu Leu Phe Cys Tyr Ser Asn Cys His
85 90 95

Lys Glu Gln Thr Ile Ala Ser Met Asn Leu Thr Val Tyr Trp Phe Pro 100 105 110

Glu His Val Glu Leu Ala Pro Leu Pro Leu Trp Gln Pro Val Gly Glu 115 120 125

Glu Leu Asn Leu Ser Cys Leu Val Ser Gly Gly Ala Pro Arg Ala His 130 \$135\$

Val Gly Lys Gly Glu Pro Ala Lys Val Met Phe Thr Val Gln Ser Arg

Arg Glu Asp His Gly Thr Asn Phe Ser Cys Arg Trp Glu Leu Asp Leu 185 Arg Ser Gln Gly Leu Glu Leu Phe Gln Asn Thr Ser Ala Pro Arg Lys Leu Gln Thr Tyr Val Leu Pro Ser Ile Asp Pro His Leu Glu Val Pro Pro Ile Val Glu Val Gly Ser Arg Trp Pro Val Asn Cys Thr Leu Asp Gly Leu Phe Pro Ala Ser Asp Ala Lys Val Tyr Leu Val Leu Gly Asp Gln Lys Leu Glu Ser Asn Ile Thr Tyr Asp Gly Asp Ser Val Leu Ala Lys Ala Trp Met Glu Glu Asn Glu Glu Gly Thr His Ser Leu Lys Cys Ser Val Thr Leu Gly Glu Val Ser Arg Arg Thr Gln Glu Asn Val Thr Val Tyr Ser Phe Pro Leu Pro Thr Leu Thr Leu Ser Pro Pro Glu Val 310 Ser Glu Trp Thr Thr Val Thr Val Glu Cys Val Thr Arg Asp Gly Ala Val Val Lys Leu Asn Gly Thr Ser Ala Val Pro Pro Gly Pro Arg Ala Gln Leu Lys Leu Asn Ala Ser Ala Ser Asp His Arg Ser Asn Phe Ser Cys Ser Ala Ala Leu Glu Ile Ala Gly Gln Val Val His Lys His Gln Thr Leu Glu Leu His Val Leu Tyr Gly Pro Arg Leu Asp Gln Arg Asp Cys Pro Gly Asn Trp Thr Trp Gln Glu Gly Ser Glu Gln Thr Leu Lys 410 Cys Glu Ala Gln Gly Asn Pro Ile Pro Lys Leu Asn Cys Ser Arg Lys Gly Asp Gly Ala Ser Leu Pro Ile Gly Asp Leu Arg Pro Val Arg Arg Glu Val Ala Gly Thr Tyr Leu Cys Arg Ala Thr Ser Ala Arg Gly Arg Val Thr Arg Glu Val Val Leu Asn Val Leu His Gly Gln Asn Ile Leu 470 475

Asp Ile Val Ile Pro Val Val Ala Val Thr Leu Ile Leu Gly Ala Leu 485 490 495

Gly Thr Ala Gly Tyr Val Tyr Asn Tyr Gln Arg Lys Ile Gln Lys Tyr 500 505 510

Glu Leu Gln Lys Ala Arg Lys Ala Gln Glu Glu Ala Ala Leu Lys Leu 515 520 525

Asn Ala Gln Ser Thr Pro Pro 530 535

<210> 85

<211> 530

<212> PRT

<213> Ovis aries

<400> 85

Met Ala Pro Gly Ala Ala Pro Ala Ala Leu Leu Ala Leu Leu Val Leu 1 5 10 15

Leu Gly Thr Leu Leu Pro Gly Ser Gly Gly Ala Glu Ile Ser Ile His 20 25 30

Pro Pro Lys Ala Ile Ile Pro Arg Gly Gly Ser Leu Arg Val Asn Cys 35 40 45

Ser Ile Ser Cys Asp Arg Lys Thr Thr Phe Gly Leu Glu Thr Val Leu 50 55 60

Asn Lys Glu Glu Val Ser Arg Gly Pro Asn Trp Lys Val Phe Glu Leu 65 70 75 80

Ser Asp Val Glu Glu Glu Ile Asn Pro Leu Cys Tyr Ser Asn Cys His 85 90 95

Gly Glu Gln Ile Val Ala Ser Met Asn Leu Thr Ile Tyr Trp Phe Pro 100 105 110

Glu Arg Val Glu Leu Ala Pro Leu Pro Leu Trp Gln Pro Val Gly Glu 115 120 125

Glu Leu Asn Leu Ser Cys Gln Val Ser Gly Gly Gly Pro Arg His His 130 135 140

Leu Ser Met Val Leu Leu Arg Gly Glu Glu Glu Leu Asp Arg Gln Pro 145 150 155 160

Val Gly Lys Glu Glu Pro Ala Glu Val Thr Phe Met Val Gln Pro Arg 165 170 175

Arg Glu Asp His Gly Thr Ser Phe Ser Cys Arg Trp Glu Leu Asp Leu 180 185 190

Arg Ser Gln Gly Leu Glu Leu Phe Gln Asn Thr Ser Ala Pro Arg Lys 195 200 205 Leu Gln Thr Tyr Val Leu Pro Ser Thr Asp Pro His Leu Glu Ala Pro Pro Val Val Glu Val Gly Ser Arg Trp Pro Val Lys Cys Thr Leu Asp 230 Gly Leu Phe Pro Ala Ser Asp Ala Glu Val Tyr Val Gln Leu Gly Asp Gln Lys Leu Glu Ser Asn Ile Thr Tyr Asn Gly Asp Ser Val Leu Ala 265 Glu Ala Trp Thr Glu Glu Asn Glu Glu Gly Thr His Ser Leu Arg Cys Ser Val Ser Leu Gly Glu Lys Ile Arg Arg Thr Arg Gly Ser Val Thr Met Tyr Ser Phe Pro Leu Pro Thr Leu Thr Leu Ser Pro Pro Glu Val 310 Ser Glu Trp Thr Thr Val Thr Val Glu Cys Val Thr Arg Asp Gly Ala 330 325 Val Val Arg Leu Asn Gly Val Ser Ala Glu Pro Pro Gly Pro Arg Ala Gln Leu Lys Leu Asn Val Ser Ala Asp Asp His Gly Ser Asn Phe Ser Cys Ser Ala Ala Leu Lys Ile Ala Gly Gln Glu Val His Lys Ile Gln Thr Arg Glu Leu His Val Leu Tyr Gly Pro Arg Leu Asp Gln Arg Asp Cys Leu Gly Asn Trp Thr Trp Gln Glu Gly Ser Glu Gln Thr Leu Lys Cys Ala Ala Arg Gly Asn Pro Ile Pro Lys Leu Asn Cys Ser Arg Lys 425 Gly Asp Gly Ala Ser Leu Pro Ile Gly Asp Leu Arg Pro Val Thr Arg Glu Val Ala Gly Thr Tyr Leu Cys Trp Ala Thr Ser Ala Arg Gly Gly Val Thr Arg Glu Val Val Leu Asn Val Leu Tyr Gly Gln Asn Ile Leu 470 Asp Ile Val Ile Pro Val Val Ala Val Thr Leu Ile Leu Gly Thr Leu 485 Gly Thr Ala Gly Tyr Ile Tyr Asn Tyr Gln Arg Lys Ile Gln Lys Tyr 505 500

Glu Leu Gln Lys Ala Gln Lys Glu Ala Ala Leu Lys Leu Lys Ser Thr 515 520 525

Pro Pro 530

<210> 86

<211> 545

<212> PRT

<213> Rattus norvegicus

<400> 86

Met Ala Ser Thr Arg Ala Arg Pro Met Leu Pro Leu Leu Leu Val Leu 1 5 10 15

Val Ala Val Val Ile Pro Gly Pro Val Gly Ala Gln Val Ser Ile His 20 25 30

Pro Thr Glu Ala Phe Leu Pro Arg Gly Gly Ser Val Gln Val Asn Cys 35 40 45

Ser Ser Cys Glu Asp Glu Asn Leu Gly Leu Gly Leu Glu Thr Asn 50 55 60

Trp Met Lys Asp Glu Leu Ser Ser Gly His Asn Trp Lys Leu Phe Lys 65 70 75 80

Leu Ser Asp Ile Gly Glu Asp Ser Arg Pro Leu Cys Phe Glu Asn Cys 85 90 95

Gly Thr Thr Gln Ser Ser Ala Ser Ala Thr Ile Thr Val Tyr Ser Phe
100 105 110

Pro Glu Arg Val Glu Leu Asp Pro Leu Pro Ala Trp Gln Gln Val Gly 115 120 125

Lys Asn Leu Ile Leu Arg Cys Leu Val Glu Gly Gly Ala Pro Arg Thr 130 135 140

Gln Leu Ser Val Val Leu Leu Arg Gly Asn Glu Thr Leu Ser Arg Gln 145 150 155 160

Ala Val Asp Gly Asp Pro Lys Glu Ile Thr Phe Thr Val Leu Ala Ser 165 170 175

Arg Gly Asp His Gly Ala Asn Phe Ser Cys Phe Thr Glu Leu Asp Leu 180 185 190

Arg Pro Gln Gly Leu Ser Leu Phe Lys Asn Val Ser Glu Val Arg Gln 195 200 205

Leu Arg Thr Phe Asp Leu Pro Thr Arg Val Leu Lys Leu Asp Thr Pro 210 215 220

Asp Leu Leu Glu Val Gly Thr Gln Gln Lys Phe Leu Cys Ser Leu Glu 225 230 235 240

- 370 375 380
- Ala Leu Glu Val Asp Gly Lys Ser Leu Phe Lys Asn Gln Thr Leu Glu 385 390 395 400
- Leu His Val Leu Tyr Gly Pro His Leu Asp Lys Lys Asp Cys Leu Gly 405 410 415
- Asn Trp Thr Trp Gln Glu Gly Ser Gln Gln Thr Leu Thr Cys Gln Pro 420 425 430
- Gln Gly Asn Pro Ala Pro Asn Leu Thr Cys Ser Arg Lys Ala Asp Gly 435 440 445
- Val Pro Leu Pro Ile Gly Met Val Lys Ser Val Lys Arg Glu Met Asn 450 455 460
- Gly Thr Tyr Lys Cys Arg Ala Phe Ser Ser Arg Gly Ser Ile Thr Arg 465 470 475 480
- Asp Val His Leu Thr Val Leu Tyr His Asp Gln Asn Thr Trp Val Ile 485 490 495
- Ile Val Gly Val Leu Val Leu Ile Ile Ala Gly Phe Val Ile Val Ala 500 505 510
- Ser Ile Tyr Thr Tyr Tyr Arg Gln Arg Lys Ile Arg Ile Tyr Lys Leu 515 520 525
- Gln Lys Ala Gln Glu Glu Ala Leu Lys Leu Lys Val Gln Ala Pro Pro 530 535 540

Pro 545

<210> 87

<211> 917

<212> PRT

<213> Rattus norvegicus

<400> 87

Met Pro Gly Pro Ser Pro Gly Leu Arg Arg Thr Leu Leu Gly Leu Trp

1 10 15

Ala Ala Leu Gly Leu Gly Ile Leu Gly Ile Ser Ala Val Ala Leu Glu 20 25 30

Pro Phe Trp Ala Asp Leu Gln Pro Arg Val Ala Leu Val Glu Arg Gly 35 40 45

Gly Ser Leu Trp Leu Asn Cys Ser Thr Asn Cys Pro Arg Pro Glu Arg
50 55 60

Gly Gly Leu Glu Thr Ser Leu Arg Arg Asn Gly Thr Gln Arg Gly Leu
65 70 75 80

Arg Trp Leu Ala Arg Gln Leu Val Asp Ile Arg Glu Pro Glu Thr Gln 85 90 95

Pro Val Cys Phe Phe Arg Cys Ala Arg Arg Thr Leu Gln Ala Arg Gly
100 105 110

Leu Ile Arg Thr Phe Gln Arg Pro Asp Arg Val Glu Leu Val Pro Leu 115 120 125

Pro Pro Trp Gln Pro Val Gly Glu Asn Phe Thr Leu Ser Cys Arg Val 130 135 140

Pro Gly Ala Gly Pro Arg Ala Ser Leu Thr Leu Thr Leu Leu Arg Gly 145 150 155 160

Gly Gln Glu Leu Ile Arg Arg Ser Phe Val Gly Glu Pro Pro Arg Ala 165 170 175

Arg Gly Ala Met Leu Thr Ala Thr Val Leu Ala Arg Arg Glu Asp His 180 185 190

Arg Ala Asn Phe Ser Cys Leu Ala Glu Leu Asp Leu Arg Pro His Gly
195 200 205

Leu Gly Leu Phe Ala Asn Ser Ser Ala Pro Arg Gln Leu Arg Thr Phe 210 215 220

Ala Met Pro Pro Leu Ser Pro Ser Leu Ile Ala Pro Arg Phe Leu Glu 225 230 235 . 240

Val Gly Ser Glu Arg Pro Val Thr Cys Thr Leu Asp Gly Leu Phe Pro 245 250 255

- Ala Pro Glu Ala Gly Val Tyr Leu Ser Leu Gly Asp Gln Arg Leu His 260 265 270
- Pro Asn Val Thr Leu Asp Gly Glu Ser Leu Val Ala Thr Ala Thr Ala 275 280 285
- Thr Ala Ser Glu Glu Glu Gly Thr Lys Gln Leu Met Cys Ile Val 290 295 300
- Thr Leu Gly Gly Glu Ser Arg Glu Thr Gln Glu Asn Leu Thr Val Tyr 305 310 315 320
- Ser Phe Pro Ala Pro Leu Leu Thr Leu Ser Glu Pro Glu Ala Pro Glu 325 330 335
- Gly Lys Met Val Thr Val Ser Cys Trp Ala Gly Ala Arg Ala Leu Val 340 345 350
- Thr Leu Glu Gly Ile Pro Ala Ala Val Pro Gly Gln Pro Ala Glu Leu 355 360 365
- Gln Leu Asn Val Thr Lys Asn Asp Asp Lys Arg Gly Phe Phe Cys Asp 370 375 380
- Ala Ala Leu Asp Val Asp Gly Glu Thr Leu Arg Lys Asn Gln Ser Ser 385 390 395 400
- Glu Leu Arg Val Leu Tyr Ala Pro Arg Leu Asp Asp Leu Asp Cys Pro
  405 410 415
- Arg Ser Trp Thr Trp Pro Glu Gly Pro Glu Gln Thr Leu His Cys Glu 420 425 430
- Ala Arg Gly Asn Pro Glu Pro Ser Val His Cys Ala Arg Pro Asp Gly 435 440 445
- Gly Ala Val Leu Ala Leu Gly Leu Leu Gly Pro Val Thr Arg Ala Leu 450 455 460
- Ala Gly Thr Tyr Arg Cys Thr Ala Ile Asn Gly Gln Gly Gln Ala Val 465 470 475 480
- Lys Asp Val Thr Leu Thr Val Glu Tyr Ala Pro Ala Leu Asp Ser Val 485 490 490
- Gly Cys Pro Glu Arg Ile Thr Trp Leu Glu Gly Thr Glu Ala Ser Leu 500 510
- Ser Cys Val Ala His Gly Val Pro Pro Pro Ser Val Ser Cys Val Arg 515 520 525
- Ser Gly Lys Glu Glu Val Met Glu Gly Pro Leu Arg Val Ala Arg Glu 530 535 540
- His Ala Gly Thr Tyr Arg Cys Glu Ala Ile Asn Ala Arg Gly Ser Ala 545 550 555 560

Ala Lys Asn Val Ala Val Thr Val Glu Tyr Gly Pro Ser Phe Glu Glu 565 Leu Gly Cys Pro Ser Asn Trp Thr Trp Val Glu Gly Ser Gly Lys Leu 585 Phe Ser Cys Glu Val Asp Gly Lys Pro Glu Pro Arg Val Glu Cys Val Gly Ser Glu Gly Ala Ser Glu Gly Val Val Leu Pro Leu Val Ser Ser 615 Asn Ser Gly Ser Arg Asn Ser Met Thr Pro Gly Asn Leu Ser Pro Gly Ile Tyr Leu Cys Asn Ala Thr Asn Arg His Gly Ser Thr Val Lys Thr Val Val Val Ser Ala Glu Ser Pro Pro Gln Met Asp Glu Ser Ser Cys Pro Ser His Gln Thr Trp Leu Glu Gly Ala Glu Ala Thr Ala Leu Ala 680 Cys Ser Ala Arg Gly Arg Pro Ser Pro Arg Val Arg Cys Ser Arg Glu 695 Gly Ala Ala Arg Leu Glu Arg Leu Gln Val Ser Arg Glu Asp Ala Gly Thr Tyr Leu Cys Val Ala Thr Asn Ala His Gly Thr Asp Ser Arg Thr Val Thr Val Gly Val Glu Tyr Arg Pro Val Val Ala Glu Leu Ala Ala Ser Pro Pro Ser Val Arg Pro Gly Gly Asn Phe Thr Leu Thr Cys Arg Ala Glu Ala Trp Pro Pro Ala Gln Ile Ser Trp Arg Ala Pro Pro Gly Ala Leu Asn Leu Gly Leu Ser Ser Asn Asn Ser Thr Leu Ser Val Ala Gly Ala Met Gly Ser His Gly Gly Glu Tyr Glu Cys Ala Ala Thr Asn 805 Ala His Gly Arg His Ala Arg Arg Ile Thr Val Arg Val Ala Gly Pro 825 Trp Leu Trp Val Ala Val Gly Gly Ala Ala Gly Gly Ala Ala Leu Leu Ala Ala Gly Ala Gly Leu Ala Phe Tyr Val Gln Ser Thr Ala Cys Lys

855

Lys Gly Glu Tyr Asn Val Gln Glu Ala Glu Ser Ser Gly Glu Ala Val 865 870 875 880

Cys Leu Asn Gly Ala Gly Gly Thr Pro Gly Ala Glu Gly Gly Ala Glu 885 890 895

Thr Pro Gly Thr Ala Glu Ser Pro Ala Asp Gly Glu Val Phe Ala Ile 900 905 910

Gln Leu Thr Ser Ser 915

<210> 88

<211> 151

<212> PRT

<213> Homo sapiens

<220>

<221> MOD\_RES

<222> (12)

<223> Any amino acid

<220>

<221> MOD RES

<222> (77)..(81)

<223> Any amino acid

<220>

<221> MOD\_RES

<222> (132)

<223> Any amino acid

<220>

<221> MOD\_RES

<222> (145)..(147)

<223> Any amino acid

<400> 88

Glu Asp Ser Gln Pro Met Cys Tyr Ser Asn Cys Xaa Asp Gly Gln Ser 1 5 10 15

Thr Ala Lys Thr Phe Leu Thr Val Tyr Trp Thr Pro Glu Arg Val Glu 20 25 30

Leu Ala Pro Leu Pro Ser Trp Gln Pro Val Gly Lys Asn Leu Thr Leu 35 40 45

Arg Cys Gln Val Glu Gly Gly Ala Pro Arg Ala Asn Leu Thr Val Val 50 55 60

Leu Leu Arg Gly Glu Lys Glu Leu Lys Arg Glu Pro Xaa Xaa Xaa 65 70 75 80

Xaa Ala Glu Val Thr Thr Thr Val Leu Val Arg Arg Asp His His Gly 85 90 95

Ala Asn Phe Ser Cys Arg Thr Glu Leu Asp Leu Arg Pro Gln Gly Leu 100 105 110

Glu Leu Phe Glu Asn Thr Ser Ala Pro Tyr Gln Leu Gln Thr Phe Val 115 120 125

Leu Pro Ala Xaa Pro Pro Gln Leu Val Ser Pro Arg Val Leu Glu Val 130 135 140

<210> 89

<211> 1252

<212> PRT

<213> Rattus norvegicus

<400> 89

Met Gly Ala Lys Arg Val Thr Val Arg Gly Ala Arg Thr Ser Pro Ile
1 5 10 15

His Arg Met Ser Ser Leu Thr Pro Leu Leu Met Gly Met Leu Thr 20 25 30

Ser Gly Leu Ala Glu Ser Pro Val Pro Thr Ser Ala Pro Arg Gly Phe 35 40 45

Trp Ala Leu Ser Glu Asn Leu Thr Ala Val Glu Gly Thr Thr Val Lys
50 55 60

Leu Trp Cys Gly Val Arg Ala Pro Gly Ser Val Val Gln Trp Ala Lys
65 70 75 80

Asp Gly Leu Leu Gly Pro Asn Pro Lys Met Pro Gly Phe Pro Arg 85 90 95

Tyr Ser Leu Glu Gly Asp Arg Ala Lys Gly Glu Phe His Leu Leu Ile 100 105 110

Glu Ala Cys Asp Leu Ser Asp Asp Ala Glu Tyr Glu Cys Gln Val Gly

Arg Ser Glu Leu Gly Pro Glu Leu Val Ser Pro Lys Val Ile Leu Ser 130 135 140

Thr Val Thr Trp Val Ala Gly Gln Glu Tyr Val Val Thr Cys Val Ser 165 170 175

Gly Asp Ala Lys Pro Ala Pro Asp Ile Thr Phe Ile Gln Ser Gly Arg 180 185 190

Thr Ile Leu Asp Val Ser Ser Asn Val Asn Glu Gly Ser Glu Glu Lys 195 200 205 Leu Cys Ile Thr Glu Ala Glu Ala Arg Val Ile Pro Gln Ser Ser Asp 215 Asn Gly Gln Leu Leu Val Cys Glu Gly Ser Asn Pro Ala Leu Asp Thr 235 230 Pro Ile Lys Ala Ser Phe Thr Met Asn Ile Leu Phe Pro Pro Gly Pro Pro Val Ile Asp Trp Pro Gly Leu Asn Glu Gly His Val Arg Ala Gly 265 Glu Asn Leu Glu Leu Pro Cys Thr Ala Arg Gly Gly Asn Pro Pro Ala 275 Thr Leu Gln Trp Leu Lys Asn Gly Lys Pro Val Ser Thr Ala Trp Gly 295 Thr Glu His Ala Gln Ala Val Ala His Ser Val Leu Val Met Thr Val 310 Arg Pro Glu Asp His Gly Ala Arg Leu Ser Cys Gln Ser Tyr Asn Ser Val Ser Ala Gly Thr Gln Glu Arg Ser Ile Thr Leu Gln Val Thr Phe 345 Pro Pro Ser Ala Ile Thr Ile Leu Gly Ser Val Ser Gln Ser Glu Asn Lys Asn Val Thr Leu Cys Cys Leu Thr Lys Ser Ser Arg Pro Arg Val Leu Leu Arg Trp Trp Leu Gly Gly Arg Gln Leu Leu Pro Thr Asp Glu Thr Val Met Asp Gly Leu His Gly Gly His Ile Ser Met Ser Asn Leu Thr Phe Leu Val Arg Arg Glu Asp Asn Gly Leu Pro Leu Thr Cys Glu 420 Ala Phe Ser Asp Ala Phe Ser Lys Glu Thr Phe Lys Lys Ser Leu Thr 440 Leu Asn Val Lys Tyr Pro Ala Gln Lys Leu Trp Ile Glu Gly Pro Pro 455 450 Glu Gly Gln Tyr Ile Arg Thr Gly Thr Arg Val Arg Leu Val Cys Leu 475 470 Ala Ile Gly Gly Asn Pro Asp Pro Ser Leu Ile Trp Phe Lys Asp Ser 485 490 Arg Pro Val Ser Glu Pro Arg Gln Pro Gln Glu Pro Arg Arg Val Gln 505

- Leu Gly Ser Val Glu Lys Ser Gly Ser Thr Phe Ser Arg Glu Leu Val 515 520 525
- Leu Ile Ile Gly Pro Pro Asp Asn Arg Ala Lys Phe Ser Cys Lys Ala 530 535 540
- Gly Gln Leu Ser Ala Ser Thr Gln Leu Val Val Gln Phe Pro Pro Thr 545 550 555 560
- Asn Leu Thr Ile Leu Ala Asn Ser Ser Ala Leu Arg Pro Gly Asp Ala 565 570 575
- Leu Asn Leu Thr Cys Val Ser Ile Ser Ser Asn Pro Pro Val Asn Leu 580 585 590
- Ser Trp Asp Lys Glu Gly Glu Arg Leu Glu Asp Val Ala Ala Lys Pro 595 600 605
- Gln Ser Ala Pro Phe Lys Gly Ser Ala Ala Ser Arg Ser Val Phe Leu 610 615 620
- Arg Val Ser Ser Arg Asp His Gly Gln Arg Val Thr Cys Arg Ala His 625 630 635 640
- Ser Glu Ala Leu Arg Glu Thr Val Ser Ser Phe Tyr Arg Phe Asn Val 645 650 655
- Leu Tyr Pro Pro Glu Phe Leu Gly Glu Gln Val Arg Ala Val Thr Val 660 665 670
- Val Glu Gln Gly Gln Val Leu Leu Pro Val Ser Val Ser Ala Asn Pro 675 680 685
- Ala Pro Glu Ala Phe Asn Trp Thr Phe Arg Gly Tyr Arg Leu Ser Pro 690 695 700
- Ala Gly Gly Pro Arg His Arg Ile Leu Ser Gly Gly Ala Leu Gln Leu 705 710 715 720
- Trp Asn Val Thr Arg Ala Asp Asp Gly Phe Tyr Gln Leu His Cys Gln 725 730 735
- Asn Ser Glu Gly Thr Ala Glu Ala Leu Leu Lys Leu Asp Val His Tyr 740 745 750
- Ala Pro Thr Ile Arg Ala Leu Arg Asp Pro Thr Glu Val Asn Val Gly 755 760 765
- Gly Ser Val Asp Ile Val Cys Thr Val Asp Ala Asn Pro Ile Leu Pro 770 775 780
- Glu Met Phe Ser Trp Glu Arg Leu Gly Glu Glu Glu Glu Asp Leu Asn 785 790 795 800
- Leu Asp Asp Met Glu Lys Val Ser Lys Gly Ser Thr Gly Arg Leu Arg 805 810 815

- Ile Arg Gln Ala Lys Leu Ser Gln Ala Gly Ala Tyr Gln Cys Ile Val 820 825 830
- Asp Asn Gly Val Ala Pro Ala Ala Arg Gly Leu Val Arg Leu Val Val 835 840 845
- Arg Phe Ala Pro Gln Val Asp Gln Pro Thr Pro Leu Thr Lys Val Ala 850 860
- Ala Ala Gly Asp Ser Thr Ser Ser Ala Thr Leu His Cys Arg Ala Arg 865 870 875 880
- Gly Val Pro Asn Ile Asp Phe Thr Trp Thr Lys Asn Gly Val Pro Leu 885 890 895
- Asp Leu Gln Asp Pro Arg Tyr Thr Glu His Arg Tyr His Gln Gly Val 900 905 910
- Val His Ser Ser Leu Leu Thr Ile Ala Asn Val Ser Ala Ala Gln Asp 915 920 925
- Tyr Ala Leu Phe Lys Cys Thr Ala Thr Asn Ala Leu Gly Ser Asp His 930 940
- Thr Asn Ile Gln Leu Val Ser Ile Ser Arg Pro Asp Pro Pro Leu Gly 945 950 955 960
- Leu Lys Val Val Ser Ile Ser Pro His Ser Val Gly Leu Glu Trp Lys 965 970 975
- Pro Gly Phe Asp Gly Gly Leu Pro Gln Arg Phe Gln Ile Arg Tyr Glu 980 985 990
- Ala Leu Glu Thr Pro Gly Phe Leu His Val Asp Val Leu Pro Thr Gln 995 1000 1005
- Ala Thr Thr Phe Thr Leu Thr Gly Leu Lys Pro Ser Thr Arg Tyr Arg 1010 1015 1020
- Ile Trp Leu Leu Ala Ser Asn Ala Leu Gly Asp Ser Gly Leu Thr Asp 1025 1030 1035 1040
- Lys Gly Ile Gln Val Ser Val Thr Thr Pro Gly Pro Asp Gln Ala Pro 1045 1050 1055
- Glu Asp Thr Asp His Gln Leu Pro Thr Glu Leu Pro Pro Gly Pro Pro 1060 1065 1070
- Arg Leu Pro Leu Leu Pro Val Leu Phe Ala Val Gly Gly Leu Leu Leu 1075 1080 1085
- Leu Ser Asn Ala Ser Cys Val Gly Gly Leu Leu Trp Arg Arg Leu 1090 1095 1100
- Arg Arg Leu Ala Glu Glu Ile Ser Glu Lys Thr Glu Ala Gly Ser Glu 1105 1115 1120

Asp Arg Ile Arg Asn Glu Tyr Glu Glu Ser Gln Trp Thr Gly Asp Arg 1125 1130 1135

Asp Thr Arg Ser Ser Thr Val Ser Thr Ala Glu Val Asp Pro Asn Tyr 1140 1145 1150

Tyr Ser Met Arg Asp Phe Ser Pro Gln Leu Pro Pro Thr Leu Glu Glu 1155 1160 1165

Val Leu Tyr His Gln Gly Ala Glu Gly Glu Asp Met Ala Phe Pro Gly 1170 1175 1180

His Leu His Asp Glu Val Glu Arg Ala Tyr Gly Pro Pro Gly Ala Trp 1185 1190 1195 1200

Gly Pro Leu Tyr Asp Glu Val Arg Met Asp Pro Tyr Asp Leu Arg Trp 1205 1210 1215

Pro Glu Val Gln Cys Glu Asp Pro Arg Gly Ile Tyr Asp Gln Val Ala 1220 1225 1230

Ala Asp Met Asp Ala Val Glu Ala Ser Ser Leu Pro Phe Glu Leu Arg 1235 1240 1245

Gly His Leu Val 1250

<210> 90

<211> 1256

<212> PRT

<213> Mus musculus

<400> 90

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His Arg Thr Cys Arg Leu Ile Pro Leu Leu Leu Ala Gly Met Leu Thr 20 25 30

Thr Gly Leu Ala Gln Ser Pro Val Pro Thr Ser Ala Pro Arg Gly Phe 35 40 45

Trp Ala Leu Ser Glu Asn Leu Thr Val Val Glu Gly Ser Thr Ile Lys
50 55 60

Leu Trp Cys Gly Val Arg Ala Pro Gly Ser Val Val Gln Trp Ala Lys 65 70 75 80

Asp Gly Leu Leu Gly Pro Asn Pro Lys Ile Pro Gly Phe Pro Arg

Tyr Ser Leu Glu Gly Asp Ser Ala Lys Gly Glu Phe His Leu Leu Ile 100 105 110

Glu Ala Cys Asp Leu Ser Asp Asp Ala Glu Tyr Glu Cys Gln Val Gly
115 120 125

Arg	Ser 130	Glu	Leu	Gly	Pro	Glu 135	Leu	Val	Ser	Pro	Arg 140	Val	Ile	Leu	Ser
Val 145	Leu	Val	Pro	Pro	Lys 150	Val	Leu	Gln	Leu	Thr 155	Pro	Glu	Ala	Gly	Ser 160
Thr	Val	Thr	Trp	Val 165	Ala	Gly	Gln	Glu	Tyr 170	Val	Val	Thr	Cys	Val 175	Ser
Gly	Gly	Ala	Lys 180	Pro	Ala	Pro	Asp	Ile 185	Ile	Phe	Ile	Gln	Gly 190	Gly	Arg
Thr	Val	Glu 195	Asp	Val	Ser	Ser	Ser 200	Val	Asn	Glu	Gly	Ser 205	Glu	Glu	Lys
Leu	Phe 210	Phe	Thr	Glu	Ala	Glu 215	Ala	Arg	Val	Thr	Pro 220	Gln	Ser	Ser	Asp
Asn 225	Gly	Gln	Leu	Leu	Val 230	Cys	Glu	Gly	Ser	Asn 235	Pro	Ala	Leu	Ala	Thr 240
Pro	Ile	Lys	Ala	Ser 245	Phe	Thr	Met	Asn	Ile 250	Leu	Phe	Pro	Pro	Gly 255	Pro
Pro	Val	Ile	Asp 260	Trp	Pro	Gly	Leu	Asn 265	Glu	Gly	His	Val	Arg 270	Ala	Gly
Glu	Asn	Leu 275	Glu	Leu	Pro	Cys	Ile 280	Ala	Arg	Gly	Gly	Asn 285	Pro	Pro	Ala
Thr	Leu 290	Gln	Trp	Leu	Lys	Asn 295	Gly	Lys	Pro	Val	Ser 300	Ile	Ala	Trp	Gly
Thr 305	Glu	His	Ala	Gln	Ala 310	Val	Ala	His	Ser	Val 315	Leu	Val	Met	Thr	Val 320
Arg	Pro	Glu	Asp	His 325	Gly	Ala	Arg	Leu	Ser 330	Cys	Gln	Ser	Tyr	Asn 335	Ser
Val	Ser	Ala	Glu 340	Thr	Gln	Glu	Arg	Ser 345	Ile	Thr	Leu	Gln	Val 350	Thr	Phe
Pro	Pro	Ser 355	Ala	Val	Thr	Ile	Leu 360	Gly	Ser	Thr	Ser	Gln 365	Ser	Glu	Asn
Lys	Asn 370	Val	Thr	Leu	Cys	Cys 375	Leu	Thr	Lys	Ser	Ser 380	Arg	Pro	Arg	Val
Leu 385	Leu	Arg	Trp	Trp	Leu 390	Gly	Gly	Arg	Gln	Leu 395	Leu	Pro	Thr	Asp	Glu 400
Thr	Val	Met	Asp	Gly 405	Leu	His	Gly	Gly	His 410	Ile	Ser	Met	Ser	Asn 415	Leu
Thr	Leu	Leu	Val 420	Lys	Arg	Glu	Asp	Asn 425	Gly	Leu	Ser	Leu	Thr 430	Cys	Glu

Ala Phe Ser Asp Ala Phe Ser Lys Glu Thr Phe Lys Lys Ser Leu Thr 440 Leu Asn Val Lys Tyr Pro Ala Gln Lys Leu Trp Ile Glu Gly Pro Pro Glu Gly Gln Ser Ile Arg Thr Gly Thr Arg Val Arg Leu Val Cys Leu Ala Ile Gly Gly Asn Pro Glu Pro Ser Leu Thr Trp Leu Lys Asp Ser 490 Arg Pro Val Asn Asp Pro Arg Gln Ser Gln Glu Pro Arg Arg Val Gln Leu Gly Ser Val Glu Lys Ser Gly Ser Thr Phe Ser Arg Glu Leu Val 520 Leu Ile Ile Gly Pro Pro Asp Asn Leu Ala Lys Phe Ser Cys Lys Ala Gly Gln Leu Ser Ala Ser Thr Gln Leu Val Val Gln Phe Pro Pro Thr Asn Leu Thr Ile Leu Ala Asn Ser Ser Ala Leu Arg Pro Gly Asp Ala 570 565 Leu Asn Leu Thr Cys Val Ser Ile Ser Ser Asn Pro Pro Val Asn Leu 585 Ser Leu Asp Lys Glu Gly Glu Arg Leu Asp Asp Val Ala Ala Lys Pro Gln Ser Ala Pro Phe Lys Gly Ser Ala Ala Ser Arg Ser Val Phe Leu Arg Val Ser Ser Arg Asp His Gly His Arg Val Thr Cys Arg Ala His 625 Ser Glu Ala Leu Arg Glu Thr Val Ser Ser Phe Tyr Arg Leu Asn Val Leu Tyr Pro Pro Glu Phe Leu Gly Glu Gln Val Arg Ala Val Thr Val 665 Val Glu Gln Gly Gln Ala Leu Leu Pro Val Ser Val Ser Ala Asn Pro Ala Pro Glu Ala Phe Asn Trp Thr Phe Arg Gly Tyr Arg Leu Ser Pro Ala Gly Gly Pro Arg His Arg Ile Leu Ser Gly Gly Ala Leu Gln Leu Trp Asn Val Thr Arg Ala Asp Asp Gly Phe Tyr Gln Leu His Cys Gln 730

725

Asn Ser Glu Gly Thr Ala Glu Ala Leu Leu Lys Leu Asp Val His Tyr 740 745 750

Ala Pro Thr Ile Arg Ala Leu Lys Asp Pro Thr Glu Val Asn Val Gly 755 760 765

Gly Ser Val Asp Ile Val Cys Thr Val Asp Ala Asn Pro Ile Leu Pro 770 780

Glu Met Phe Ser Trp Glu Arg Leu Gly Glu Asp Glu Glu Glu Leu Asn 785 790 795 800

Leu Asp Asp Met Glu Lys Met Ser Lys Gly Ser Thr Gly Arg Leu Arg 805 810 815

Ile Arg Gln Ala Lys Leu Ser Gln Ala Gly Ala Tyr Gln Cys Ile Val 820 825 830

Asp Asn Gly Val Ala Pro Ala Ala Arg Gly Leu Val Arg Leu Val Val 835 840 845

Arg Phe Ala Pro Gln Val Asp His Pro Thr Pro Leu Thr Lys Val Ala 850 860

Ala Ala Gly Asp Ser Thr Ser Ser Ala Thr Leu His Cys Arg Ala Arg 865 870 875 880

Gly Val Pro Asn Ile Asp Phe Thr Trp Thr Lys Asn Gly Val Pro Leu 885 890 895

Asp Leu Gln Asp Pro Arg Tyr Thr Glu His Lys Tyr His Gln Gly Val 900 905 910

Val His Ser Ser Leu Leu Thr Ile Ala Asn Val Ser Ala Ala Gln Asp 915 920 925

Tyr Ala Leu Phe Lys Cys Thr Ala Thr Asn Ala Leu Gly Ser Asp His 930 935 940

Thr Asn Ile Gln Leu Val Ser Ile Ser Arg Pro Asp Pro Pro Leu Gly 945 950 955 960

Leu Lys Val Val Ser Val Ser Pro His Ser Val Gly Leu Glu Trp Lys 965 970 975

Pro Gly Phe Asp Gly Gly Leu Pro Gln Arg Phe Gln Ile Arg Tyr Glu 980 985 990

Ala Leu Glu Thr Pro Gly Phe Leu Tyr Met Asp Val Leu Pro Ala Gln 995 1000 1005

Ala Thr Thr Phe Thr Leu Thr Gly Leu Lys Pro Ser Thr Arg Tyr Arg 1010 1015 1020

Ile Trp Leu Leu Ala Ser Asn Ala Leu Gly Asp Ser Gly Leu Thr Asp 1025 1030 1035 1040 Lys Gly Ile Gln Val Ser Ile Thr Thr Pro Gly Leu Asp Gln Ala Pro 1045 1050 1055

Glu Asp Thr Asp Gln Pro Leu Pro Thr Glu Gln Pro Pro Gly Pro Pro 1060 1065 1070

Arg Leu Pro Leu Leu Pro Val Leu Phe Ala Val Gly Gly Leu Leu Leu 1075 1080 1085

Leu Ser Asn Ala Ser Cys Val Gly Gly Leu Leu Trp Arg Arg Leu 1090 1095 1100

Arg Arg Leu Ala Glu Glu Ile Ser Glu Lys Thr Glu Ala Gly Ser Glu 1105 1115 1120

Glu Asp Arg Ile Arg Asn Glu Tyr Glu Glu Ser Gln Trp Thr Gly Asp 1125 1130 1135

Arg Asp Thr Arg Ser Ser Thr Val Ser Thr Ala Glu Val Asp Pro His
1140 1145 1150

Tyr Tyr Ser Met Arg Asp Phe Ser Pro Gln Leu Pro Pro Thr Leu Glu 1155 1160 1165

Glu Val Ser Tyr Arg Gln Ala Phe Thr Gly Ile Glu Asp Glu Asp Met 1170 1175 1180

Ala Phe Pro Gly His Leu Tyr Asp Glu Val Glu Arg Val Tyr Gly Pro 1185 1190 1195 1200

Pro Gly Val Trp Gly Pro Leu Tyr Asp Glu Val Gln Met Asp Pro Tyr 1205 1210 1215

Asp Leu Arg Trp Pro Glu Val Lys Tyr Glu Asp Pro Arg Gly Ile Tyr 1220 1225 1230

Asp Gln Val Ala Ala Asp Met Asp Ala Gly Glu Pro Gly Ser Leu Pro 1235 1240 1245

Phe Glu Leu Arg Gly His Leu Val 1250 1255

<210> 91

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 91

gacctgtact tcattttgga caaatcagg

29

<210> 92

<211> 34

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<220> <223> I	Description of Artificial Sequence: Primer	
<400> 9 gagctca		34
<210> 9 <211> 2 <212> I <213> I	27	
<220> <223> I	Description of Artificial Sequence: Primer	
<400> s	• -	27
<210> 3 <211> 4 <212> 1 <213> 2	49	
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<400> gtccaa		49
<210><211><211><212><213>	28	
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<400> ggggga	95 legea gggaggatgg gggteeag	28
<210><211><212><213>	30	
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<400> gagcto	96 eccgt cagaacagtg tgtggtgg	30

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<210> 97
<211> 46
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 97
                                                                   46
atcctccctg cgtcccccag ccaaactagt agaggtgaac aaaagc
<210> 98
<211> 1650
<212> DNA
<213> Unknown Organism
<220>
<221> CDS
<222> (34)..(1647)
<223> Description of Unknown Organism: ATR-IgA2 fusion
      nucleotide
<400> 98
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                                      Met Ala Ser Lys Pro Phe Leu
tct ctt ctt tct ttg tct ttg ctt ttg ttc acc tct act agt ttg gct
                                                                   102
Ser Leu Leu Ser Leu Ser Leu Leu Leu Phe Thr Ser Thr Ser Leu Ala
                             15
         10
gac ctg tac ttc att ttg gac aaa tca gga agt gtg ctg cac cac tgg
                                                                   150
Asp Leu Tyr Phe Ile Leu Asp Lys Ser Gly Ser Val Leu His His Trp
                         30
aat gaa atc tat tac ttt gtg gaa cag ttg gct cac aaa ttc atc agc
                                                                   198
Asn Glu Ile Tyr Tyr Phe Val Glu Gln Leu Ala His Lys Phe Ile Ser
 40
cca cag ttg aga atg tcc ttt att gtt ttc tcc acc cga gga aca acc
                                                                   246
Pro Gln Leu Arg Met Ser Phe Ile Val Phe Ser Thr Arg Gly Thr Thr
                 60
tta atg aaa ctg aca gaa gac aga gaa caa atc cgt caa ggc cta gaa
                                                                    294
Leu Met Lys Leu Thr Glu Asp Arg Glu Gln Ile Arg Gln Gly Leu Glu
             75
gaa ctc cag aaa gtt ctg cca gga gga gac act tac atg cat gaa gga
Glu Leu Gln Lys Val Leu Pro Gly Gly Asp Thr Tyr Met His Glu Gly
         90
ttt gaa agg gcc agt gag cag att tat tat gaa aac aga caa ggg tac
Phe Glu Arg Ala Ser Glu Gln Ile Tyr Tyr Glu Asn Arg Gln Gly Tyr
    105
                         110
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agg Arg 120	aca Thr	gcc Ala	agc Ser	gtc Val	atc Ile 125	att Ile	gct Ala	ttg Leu	act Thr	gat Asp 130	gga Gly	gaa Glu	ctc Leu	cat His	gaa Glu 135	438
gat Asp	ctc Leu	ttt Phe	ttc Phe	tat Tyr 140	tca Ser	gag Glu	agg Arg	gag Glu	gct Ala 145	aat Asn	agg Arg	tct Ser	cga Arg	gat Asp 150	ctt Leu	486
ggt Gly	gca Ala	att Ile	gtt Val 155	tac Tyr	tgt Cys	gtt Val	ggt Gly	gtg Val 160	aaa Lys	gat Asp	ttc Phe	aat Asn	gag Glu 165	aca Thr	cag Gln	534
ctg Leu	gcc Ala	cgg Arg 170	att Ile	gcg Ala	gac Asp	agt Ser	aag Lys 175	gat Asp	cat His	gtg Val	ttt Phe	ccc Pro 180	gtg Val	aat Asn	gac Asp	582
ggc Gly	ttt Phe 185	cag Gln	gct Ala	ctg Leu	caa Gln	ggc Gly 190	atc Ile	atc Ile	cac His	tca Ser	att Ile 195	ttg Leu	agc Ser	tct Ser	gct Ala	630
tcc Ser 200	cca Pro	acc Thr	agc Ser	cct Pro	aag Lys 205	gtc Val	ttc Phe	cct Pro	ctc Leu	agc Ser 210	ctt Leu	gac Asp	agc Ser	acc Thr	cct Pro 215	678
											cag Gln					726
											gga Gly					774
											ggt Gly					822
											tgc Cys 275					870
tcc Ser 280	gtt Val	acc Thr	tgc Cys	cat His	gtt Val 285	aag Lys	cac His	tac Tyr	acc Thr	aac Asn 290	tcc Ser	agc Ser	cag Gln	gat Asp	gtt Val 295	918
act Thr	gtt Val	cca Pro	tgc Cys	cgt Arg 300	gtt Val	cca Pro	cca Pro	cct Pro	cca Pro 305	cca Pro	tgc Cys	tgc Cys	cac His	cca Pro 310	cgt Arg	966
											ctc Leu					1014
				_						_	gat Asp	_			-	1062

acc Thr	ttc Phe 345	acc Thr	tgg Trp	acc Thr	cca Pro	agc Ser 350	tct Ser	ggt Gly	aag Lys	agc Ser	gct Ala 355	gtt Val	caa Gln	gga Gly	cca Pro	1110
cct Pro 360	gag Glu	cgt Arg	gac Asp	ctc Leu	tgt Cys 365	gga Gly	tgc Cys	tac Tyr	tct Ser	gtt Val 370	agc Ser	tct Ser	gtt Val	ctt Leu	cct Pro 375	1158
ggt Gly	tgt Cys	gcc Ala	cag Gln	cct Pro 380	tgg Trp	aac Asn	cac His	ggt Gly	gag Glu 385	acc Thr	ttc Phe	acc Thr	tgc Cys	act Thr 390	gct Ala	1206
gcc Ala	cac His	cca Pro	gag Glu 395	ttg Leu	aag Lys	acc Thr	cca Pro	ctt Leu 400	acc Thr	gcc Ala	aac Asn	atc Ile	acc Thr 405	aag Lys	tcc Ser	1254
								cac His								1302
								acc Thr								1350
								agg Arg								1398
								tgg Trp								1446
cag Gln	gga Gly	act Thr	acc Thr 475	acc Thr	tac Tyr	gct Ala	gtt Val	acc Thr 480	agc Ser	atc Ile	ctt Leu	cgt Arg	gtt Val 485	gct Ala	gct Ala	1494
gag Glu	gac Asp	tgg Trp 490	aag Lys	aag Lys	ggt Gly	gag Glu	acc Thr 495	ttc Phe	tcc Ser	tgc Cys	atg Met	gtt Val 500	ggt Gly	cac His	gag Glu	1542
								aag Lys								1590
aag Lys 520	Pro	acc Thr	cac His	atc Ile	aat Asn 525	gtt Val	tct Ser	gtt Val	gtc Val	atg Met 530	Ala	gag Glu	gct Ala	gat Asp	gga Gly 535	1638
	tgc Cys		taa													1650

<210> 99 <211> 538 <212> PRT <213> Unknown Organism

<220>

<223> Description of Unknown Organism: ATR-IgA2 fusion
 amino acid

<400> 99

Met Ala Ser Lys Pro Phe Leu Ser Leu Leu Ser Leu Ser Leu Leu Leu 1 5 10 15

Phe Thr Ser Thr Ser Leu Ala Asp Leu Tyr Phe Ile Leu Asp Lys Ser 20 25 30

Gly Ser Val Leu His His Trp Asn Glu Ile Tyr Tyr Phe Val Glu Gln 35 40

Leu Ala His Lys Phe Ile Ser Pro Gln Leu Arg Met Ser Phe Ile Val 50 55 60

Phe Ser Thr Arg Gly Thr Thr Leu Met Lys Leu Thr Glu Asp Arg Glu 65 70 75 80

Gln Ile Arg Gln Gly Leu Glu Glu Leu Gln Lys Val Leu Pro Gly Gly 85 90 95

Asp Thr Tyr Met His Glu Gly Phe Glu Arg Ala Ser Glu Gln Ile Tyr 100 105 110

Tyr Glu Asn Arg Gln Gly Tyr Arg Thr Ala Ser Val Ile Ile Ala Leu 115 120 125

Thr Asp Gly Glu Leu His Glu Asp Leu Phe Phe Tyr Ser Glu Arg Glu 130 135 140

Ala Asn Arg Ser Arg Asp Leu Gly Ala Ile Val Tyr Cys Val Gly Val 145 150 155 160

Lys Asp Phe Asn Glu Thr Gln Leu Ala Arg Ile Ala Asp Ser Lys Asp

His Val Phe Pro Val Asn Asp Gly Phe Gln Ala Leu Gln Gly Ile Ile 180 185 190

His Ser Ile Leu Ser Ser Ala Ser Pro Thr Ser Pro Lys Val Phe Pro 195 200 205

Leu Ser Leu Asp Ser Thr Pro Gln Asp Gly Asn Val Val Ala Cys 210 215 220

Leu Val Gln Gly Phe Phe Pro Gln Glu Pro Leu Ser Val Thr Trp Ser 225 230 235 240

Glu Ser Gly Gln Asn Val Thr Ala Arg Asn Phe Pro Pro Ser Gln Asp

Ala Ser Gly Asp Leu Tyr Thr Thr Ser Ser Gln Leu Thr Leu Pro Ala 260 265 270

Thr Gln Cys Pro Asp Gly Lys Ser Val Thr Cys His Val Lys His Tyr 275 280 285

Thr Asn Ser Ser Gln Asp Val Thr Val Pro Cys Arg Val Pro Pro Pro 290 295 300

Pro Pro Cys Cys His Pro Arg Leu Ser Leu His Arg Pro Ala Leu Glu 305 310 315 320

Asp Leu Leu Gly Ser Glu Ala Asn Leu Thr Cys Thr Leu Thr Gly 325 330 335

Leu Arg Asp Ala Ser Gly Ala Thr Phe Thr Trp Thr Pro Ser Ser Gly 340 345 350

Lys Ser Ala Val Gln Gly Pro Pro Glu Arg Asp Leu Cys Gly Cys Tyr 355 360 365

Ser Val Ser Ser Val Leu Pro Gly Cys Ala Gln Pro Trp Asn His Gly 370 375 380

Glu Thr Phe Thr Cys Thr Ala Ala His Pro Glu Leu Lys Thr Pro Leu 385 390 395 400

Thr Ala Asn Ile Thr Lys Ser Gly Asn Thr Phe Arg Pro Glu Val His 405 410 415

Leu Leu Pro Pro Pro Ser Glu Glu Leu Ala Leu Asn Glu Leu Val Thr 420 425 430

Leu Thr Cys Leu Ala Arg Gly Phe Ser Pro Lys Asp Val Leu Val Arg 435 440 445

Trp Leu Gln Gly Ser Gln Glu Leu Pro Arg Glu Lys Tyr Leu Thr Trp 450 455 460

Ala Ser Arg Gln Glu Pro Ser Gln Gly Thr Thr Thr Tyr Ala Val Thr 465 470 475 480

Ser Ile Leu Arg Val Ala Ala Glu Asp Trp Lys Lys Gly Glu Thr Phe 485 490 495

Ser Cys Met Val Gly His Glu Ala Leu Pro Leu Ala Phe Thr Gln Lys 500 505 510

Thr Ile Asp Arg Leu Ala Gly Lys Pro Thr His Ile Asn Val Ser Val 515 520 525

Val Met Ala Glu Ala Asp Gly Thr Cys Tyr 530 535

<210> 100

<211> 6602

<212> DNA

<213> Unknown Organism

<220>
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	catgccaacc					7129